

Serial # 081992, 914

doc: 9-10-98

=> s raffinose synthase

L1 8 RAFFINOSE SYNTHASE

=> s transgenic

L2 52105 TRANSGENIC

=> s 11 and 12

L3 1 L1 AND L2

=> d 13

L3 ANSWER 1 OF 1 CAPLUS COPYRIGHT 1998 ACS

AN 1998:217308 CAPLUS

DN 128:318794

TI Cloning of cDNA for **raffinose synthase** from cucumber, and its use for preparing raffinose or **transgenic** plants low in raffinose-type oligosaccharides

IN Oosumi, Chieko; Nozaki, Shinji; Kida, Takao

PA Ajinomoto Co., Inc., Japan

SO Jpn. Kokai Tokkyo Koho, 26 pp.

CODEN: JKXXAF

PI JP 10084973 A2 980407 Heisei

AI JP 97-111124 970428

PRAI JP 96-107682 960426

JP 96-198079 960726

DT Patent

LA Japanese

=> d 11 1-8

L1 ANSWER 1 OF 8 BIOSIS COPYRIGHT 1998 BIOSIS

AN 98:339293 BIOSIS

DN 01339293

TI Characterization and gene cloning of **raffinose synthase** from *Cucumis sativus*.

AU Ohsumi C; Nozaki J; Kida T

CS Ajinomoto Co. Inc., CRL., Kawasaki 210, Japan

SO 1998 Annual Meeting of the Japanese Society of Plant Pathologists, Tokyo, Japan, May 3-5, 1998. *Plant and Cell Physiology* 39 (SUPPL.). 1998. S131. ISSN: 0032-0781

DT Conference

LA English

L1 ANSWER 2 OF 8 BIOSIS COPYRIGHT 1998 BIOSIS

AN 98:339292 BIOSIS

DN 01339292

TI Purification of **raffinose synthase** from *Cucumis sativus* leaves.

AU Nozaki J; Ohsumi C; Kida T

CS Ajinomoto Co. Inc., CRL., Kawasaki 210, Japan

SO 1998 Annual Meeting of the Japanese Society of Plant Pathologists, Tokyo, Japan, May 3-5, 1998. *Plant and Cell Physiology* 39 (SUPPL.). 1998. S131. ISSN: 0032-0781

DT Conference
LA English

L1 ANSWER 3 OF 8 BIOSIS COPYRIGHT 1998 BIOSIS
AN 90:197835 BIOSIS
DN BA89:104506
TI **RAFFINOSE SYNTHASE AND GALACTINOL SYNTHASE IN**
DEVELOPING SEEDS AND LEAVES OF LEGUMES.
AU CASTILLO E M; DE LUMEN B O; REYES P S; DE LUMEN H Z
CS DEP. NUTRITIONAL SCI., UNIV. CALIFORNIA, BERKELEY, CALIF. 94720.
SO J AGRIC FOOD CHEM 38 (2). 1990. 351-355. CODEN: JAFCAU ISSN:
0021-8561
LA English

L1 ANSWER 4 OF 8 AGRICOLA
AN 91:43724 AGRICOLA
DN IND91018838
TI **Raffinose synthase** and galactinol synthase in
developing seeds and leaves of legumes.
AU Castillo, E.M.; Lumen, B.O. de; Reyes, P.S.; Lumen, H.Z. de
CS University of The Philippines, Los Banos, Los Banos, The Philippines
AV DNAL (381 J8223)
SO Journal of agricultural and food chemistry, Feb 1990. Vol. 38, No.
2. p. 351-355
Publisher: Washington, D.C. : American Chemical Society.
CODEN: JAFCAU; ISSN: 0021-8561
NTE Includes references.
DT Article
FS U.S. Imprints not USDA, Experiment or Extension
LA English

L1 ANSWER 5 OF 8 CAPLUS COPYRIGHT 1998 ACS
AN 1998:217308 CAPLUS
DN 128:318794
TI Cloning of cDNA for **raffinose synthase** from
cucumber, and its use for preparing raffinose or transgenic plants
low in raffinose-type oligosaccharides
IN Oosumi, Chieko; Nozaki, Shinji; Kida, Takao
PA Ajinomoto Co., Inc., Japan
SO Jpn. Kokai Tokkyo Koho, 26 pp.
CODEN: JKXXAF
PI JP 10084973 A2 980407 Heisei
AI JP 97-111124 970428
PRAI JP 96-107682 960426
JP 96-198079 960726
DT Patent
LA Japanese

L1 ANSWER 6 OF 8 CAPLUS COPYRIGHT 1998 ACS
AN 1992:55624 CAPLUS
DN 116:55624
TI Distribution and immunolocalization of stachyose synthase in *Cucumis*
melo L
AU Holthaus, Uta; Schmitz, Klaus
CS Bot. Inst., Univ. Koeln, Cologne, W-5000/41, Germany
SO Planta (1991), 185(4), 479-86
CODEN: PLANAB; ISSN: 0032-0935
DT Journal
LA English

L1 ANSWER 7 OF 8 CAPLUS COPYRIGHT 1998 ACS
AN 1990:115882 CAPLUS
DN 112:115882
TI **Raffinose synthase** and galactinol synthase in
developing seeds and leaves of legumes

AU Castillo, Eugenia M.; De Lumen, Benito O.; Reyes, Pilar S.; De Lumen, Helen Z.
CS Dep. Nutr. Sci., Univ. California, Berkeley, CA, 94720, USA
SO J. Agric. Food Chem. (1990), 38(2), 351-5
CODEN: JAFCAU; ISSN: 0021-8561
DT Journal
LA English
OS CJACS

L1 ANSWER 8 OF 8 WPIDS COPYRIGHT 1998 DERWENT INFORMATION LTD
AN 98-264858 [24] WPIDS
DNN N98-208774 DNC C98-082361
TI **Raffinose synthase** gene - useful for preparation
of raffinose in transformed plant.
DC C06 D16 P13
PA (AJIN) AJINOMOTO KK
CYC 1
PI JP 10084973 A 980407 (9824)* 26 pp C12N015-09
ADT JP 10084973 A JP 97-111124 970428
PRAI JP 96-198079 960726; JP 96-107682 960426
IC ICM C12N015-09
ICS A01H005-00; C12N009-00

=> s raffinose

L4 6845 RAFFINOSE

=> s synthase

L5 89205 SYNTHASE

=> s 14 and 15

L6 107 L4 AND L5

=> s transgenic

L7 52105 TRANSGENIC

=> s 16 and 17

L8 3 L6 AND L7

=> d 18 1-3

L8 ANSWER 1 OF 3 CAPLUS COPYRIGHT 1998 ACS
AN 1998:217308 CAPLUS
DN 128:318794
TI Cloning of cDNA for **raffinose synthase** from
cucumber, and its use for preparing **raffinose** or
transgenic plants low in **raffinose**-type
oligosaccharides
IN Oosumi, Chieko; Nozaki, Shinji; Kida, Takao
PA Ajinomoto Co., Inc., Japan
SO Jpn. Kokai Tokkyo Koho, 26 pp.
CODEN: JKXXAF
PI JP 10084973 A2 980407 Heisei
AI JP 97-111124 970428
PRAI JP 96-107682 960426
JP 96-198079 960726
DT Patent
LA Japanese

L8 ANSWER 2 OF 3 CAPLUS COPYRIGHT 1998 ACS

AN 1993:553371 CAPLUS
DN 119:153371
TI Nucleotide sequences for galactinol **synthase** from zucchini
and soybean
IN Kerr, Phillip S.; Pearlstein, Richard W.; Schweiger, Bruce J.;
Becker-Manley, Mary F.; Pierce, John W.
PA du Pont de Nemours, E. I., and Co., USA
SO PCT Int. Appl., 79 pp.
CODEN: PIXXD2
PI WO 9302196 A1 930204
DS W: AU, BB, BG, BR, CA, CS, FI, HU, JP, KP, KR, LK, MG, MN, MW, NO,
PL, RO, RU, SD, US
RW: AT, BE, BF, BJ, CF, CG, CH, CI, CM, DE, DK, ES, FR, GA, GB, GR,
IT, LU, MC, ML, MR, NL, SE, SN, TD, TG
AI WO 92-US6057 920724
PRAI US 91-735066 910724
DT Patent
LA English

L8 ANSWER 3 OF 3 WPIDS COPYRIGHT 1998 DERWENT INFORMATION LTD
AN 93-058793 [07] WPIDS
DNN N93-044772 DNC C93-026307
TI Nucleotide sequence of galactinol **synthase** from zucchini
and soybean - used to produce plants having altered levels of
raffinose saccharide(s) and/or sucrose.
DC C06 D16 P13
IN BECKER-MANLEY, M F; KERR, P S; PEARLSTEIN, R W; PIERCE, J W;
SCHWEIGER, B J
PA (DUPO) DU PONT DE NEMOURS & CO E I
CYC 37
PI WO 9302196 A1 930204 (9307)* EN 80 pp C12N015-54
RW: AT BE CH DE DK ES FR GB GR IT LU MC NL OA SE
W: AU BB BG BR CA CS FI HU JP KP KR LK MG MN MW NO PL RO RU SD
US
AU 9224205 A 930223 (9324) C12N015-54
ZA 9205592 A 940330 (9417) 76 pp C07H000-00
EP 604458 A1 940706 (9426) EN C12N015-54
R: FR
US 5648210 A 970715 (9734) 24 pp C12Q001-68
US 5773699 A 980630 (9833) A01H001-04
ADT WO 9302196 A1 WO 92-US6057 920724; AU 9224205 A AU 92-24205 920724;
ZA 9205592 A ZA 92-5592 920724; EP 604458 A1 EP 92-917170 920724, WO
92-US6057 920724; US 5648210 A WO 92-US6057 920724, US 94-182060
940823; US 5773699 A Cont of US 91-735066 910724, Cont of WO
92-US6057 920724, Cont of US 94-182060 940823, US 96-712702 960912
FDT AU 9224205 A Based on WO 9302196; EP 604458 A1 Based on WO 9302196;
US 5648210 A Based on WO 9302196; US 5773699 A Cont of US 5648210
PRAI US 91-735066 910724; US 94-182060 940823; US 96-712702 960912
IC ICM A01H001-04; C07H000-00; C12N015-54; C12Q001-68
ICS A01H005-00; C07H021-02; C07H021-04; C12N001-21; C12N005-10;
C12N015-11; C12N015-29; C12N015-82; C12P019-18; C12P019-34

=> d 18 2-3 ab

L8 ANSWER 2 OF 3 CAPLUS COPYRIGHT 1998 ACS
AB The cDNA for zucchini leaf and soybean seed galactinol
synthase (I) are cloned and expressed in a
transgenic plant or a host cell using an appropriate
regulatory sequence. The plant or plant cell transformed with the
cDNA operatively linked to a promoter, e.g., the 35S promoter, may
have a varied level of I and thus the desirable levels of
raffinose saccharides and sucrose. Nucleotide fragments
derived from the cDNA can be used for RFLP breeding of altered
levels of **raffinose** saccharides and sucrose traits in

soybeans. I may also be prep'd. in a host, e.g., *Escherichia coli*, and used for prep'd. galactinol.

L8 ANSWER 3 OF 3 WPIDS COPYRIGHT 1998 DERWENT INFORMATION LTD

AB WO 9302196 A UPAB: 931119

An isolated nucleic acid fragment (I) comprising a nucleotide sequence encoding plant galactinol **synthase** (GaS) is new.

Also claimed are: (1) a chimeric gene (II) capable of being expressed in transformed plants, comprising (I) operably linked to a suitable regulatory sequence; (2) a chimeric gene (III) capable of being expressed in transformed plants, comprising a nucleic acid fragment encoding antisense RNA complementary to plant GaS; (3) a chimeric gene (IV) capable of being expressed in transformed microorganisms, comprising (I) operably linked to a suitable regulatory sequence; (4) plants transformed with (II) and (III) respectively; (5) seeds obt'd. from the plants of (4); (6) a microorganism transformed with (IV); (7) methods for obtaining plants and plant cells contg. altered levels of **raffinose** saccharides and/or sucrose; (8) 2 methods for producing galactinol; (9) a method of RFLP breeding of altered levels of **raffinose** saccharides and sucrose traits in soybeans; (10) a method of varying the level of D-galactase contg. oligosaccharides of sucrose in plants in response to end-user requirements.

USE/ADVANTAGE - **Transgenic** plants that produce higher than normal levels of **raffinose** saccharides possess enhanced cold tolerance and in coniferous species will result in reduced post harvest needle abscission. **Transgenic** plants that produce lower than normal levels of **raffinose** saccharide are more easily digestible and have larger amts. of metabolisable energy. In sugar beets in particular, a decrease in **raffinose** saccharide content would improve sucrose crystallisation and overcome the need for expensive processing of sugar beet extracts. The enzyme can be used for prodn. of galactinol, and the nucleic acid fragments can be used as RFLP markers in soybean genetic studies and breeding programs

Dwg.0/3



Use to restrict search to that item

[Yeast](#)
[Volume 8](#)
[Issue 2](#)

VIEWING OPTIONS: [\[MEDLINE\]](#) [\[full MEDLINE\]](#) [\[related records\]](#)

IMP2, a nuclear gene controlling the mitochondrial dependence of galactose, maltose and raffinose utilization in *Saccharomyces cerevisiae*.

Donnini C, Lodi T, Ferrero I, Puglisi PP

Yeast 1992 Feb 8:2 83-93

Abstract

The IMP2 gene of *Saccharomyces cerevisiae* is involved in the nucleo-mitochondrial control of maltose, galactose and raffinose utilization as shown by the inability of *imp2* mutants to grow on these carbon sources in respiratory-deficient conditions or in the presence of ethidium bromide and erythromycin. The negative phenotype cannot be scored in the presence of inhibitors of respiration and oxidative phosphorylation, indicating that the role of the mitochondria in the utilization of the above-mentioned carbon sources in *imp2* mutants is not at the energetical level. Mutations in the IMP2 gene also confer many phenotypic alterations in respiratory-sufficient conditions, e.g. leaky phenotype on oxidizable carbon sources, sensitivity to heat shock and sporulation deficiency. The IMP2 gene has been cloned, sequenced and disrupted. The phenotype of null *imp2* mutants is indistinguishable from that of the originally isolated mutant.

MeSH

Amino Acid Sequence ; Base Sequence ; Carbohydrates ; Galactose ; Genes, Fungal ; Genetic Complementation Test ; Maltose ; Mitochondria ; Molecular Sequence Data ; Mutation ; Phenotype ; Plasmids ; Raffinose ; Saccharomyces cerevisiae ; Support, Non-U.S. Gov't ;

Author Address

Istituto di Genetica, Universita di Parma, Italy.

Secondary Source (links)

GENBANK/X61928



[Help](#) [Feedback](#) [BioMedNet Home](#) [Search](#) [Map](#)

© 1998 BioMedNet Limited. All rights reserved.
email: info@biomednet.com.

NCBI Entrez Nucleotide QUERY
BLAST Entrez ?
Other Formats: **FASTA** **Graphic**Links: **MEDLINE** **Protein** **Related Sequences**

LOCUS SCIMP2 2071 bp DNA **PLN** 18-AUG-1993
DEFINITION *S. cerevisiae* IMP2 gene.
ACCESSION X61928 S94332
NID g3829
KEYWORDS IMP2 gene.
SOURCE baker's yeast.
ORGANISM *Saccharomyces cerevisiae*
Eukaryotae; mitochondrial eukaryotes; Fungi; Ascomycota;
Hemiascomycetes; Saccharomycetales; Saccharomycetaceae;
Saccharomyces.
REFERENCE 1 (bases 1 to 2071)
AUTHORS Lodi,T.
TITLE Direct Submission
JOURNAL Submitted (04-SEP-1991) T. Lodi, Inst di Genetica, Viale delle
Scienze, I-43100 Parma, ITALY
REFERENCE 2 (bases 1 to 2071)
AUTHORS Donnini,C., Lodi,T., Ferrero,I. and Puglisi,P.P.
TITLE IMP2, a nuclear gene controlling the mitochondrial dependence of
galactose, maltose and raffinose utilization in *Saccharomyces*
cerevisiae
JOURNAL Yeast 8 (2), 83-93 (1992)
MEDLINE 92221693
FEATURES Location/Qualifiers
source 1..2071
/organism="Saccharomyces cerevisiae"
/db_xref="taxon:4932"
/chromosome="IX"
gene 697..1635
/gene="IMP2"
CDS 697..1635
/gene="IMP2"
/codon_start=1
/db_xref="PID:g3830"
/db_xref="SWISS-PROT:P32351"
/translation="MERGHGERGRSKKKRGERDSNVSSLRSRSRASSRSRVREEEFL
KWTVLRQDPSMRLRVVDVDSEEEGEGNDEDDDGDDMDEEESDEEQVSDIENDLEI
DEEFHYDLGMKVLPNFCTSINEVLDSSKPWIAKYEISIRGHENEGVSLEQLDGGYVRA
MQLLTKGAGAEAGNQRSFILYTDLSSESTYALTYLMGAAVNQGDTVYIVHWEPSKPTD
DSQMFANVARIRKHVMHLFDCVAGVLHDLHVVVLSTHPPYPKHLLNEMIHGLKPVALC
CSLSVILSTLQNFVCSVPILAVRKKLRAKRKGISE"
BASE COUNT 565 a 534 c 534 g 438 t
ORIGIN
1 ggtcgacgaa ccaaagctgt tgcccaggta atactgcaat tcaggcacca cctcatggta
61 ttccgaggga agcatttcct gcagccactg aggcaaacgt tcatccagct tgtgtatgcc
121 ctcgcggcat gctaggttac cataccttct aggaccagga agaggcggtg tctcgtcgat
181 caggtgtgca agcttgtcca acactcccat cagccccgtta accatggcga tcgacgacgc
241 agtagcgtgc gggtcggggt cagaacagtg cgatataaac gtagtgtact tgtgaagatg
301 atacttgatt ctgtgaatgg ctgtgacga ttgaaagtcc agcgttgtct gcgtatcgaa
361 gatacgttcc accggcgttag agaagggtggc gtgcggccaa tctacacgat ccagagacat
421 ttctttgctc gtttaatgct tggtcatatg tgctgtttt tgcaacttgc actttcttcc
481 tattttatct tttaaagaa gctttaaagg tccggaaaaa aggcttgatc aaacacaaga
541 gactattgaa agggtagta cccaaagaac caacaagaga aacaaccaag tacgcaatgc
601 agaagagcat attgctgact aaactgacgg aaaacaatcg aatctacaca gcatcaaaac
661 gaaacgcga ccacgggtga gttcgactcg gagcagatgg aaagggggcca cagggaaaga
721 gtcgttagca agaagaaaag aggcgaacga gactcaaacg tgtagcgtct gtcgcgggtcg
781 agaagcagg ccagtagccg cagcagagta agggaggaag agttcctcaa gtggaccgtg
841 ttgaggcagg acccctcgat gcggttggagg gtcgtggatg tggattctga agaggaaggt
901 gagggcaacg acgaagatga cgacgacggc gacggcgacg atatggacga ggaagagtcc

961 gatgaagagc aagtgagcga tatagagaac gatttagaga ttgacgagga gttccactac
1021 gatctggca tgaaagtgtt acccaacttt tgtaccagca taaatgaagt gctagactcc
1081 agcaagccct ggataccaa gtacgagatc agcatccgtg gccacgaaaa cgaaggcgtg
1141 tctctggac aactcgacgg aggctacgtc agagccatgc aactactcac caagggtgcc
1201 ggcgcagagg cggggacca aaggccttc atcctctaca cggacctgag cagcagtc
1261 acctacgccc tgacccatct catggcgcga gctgtcaacc agggagacac cgtctacatt
1321 gtccactggg agccctcgaa gcccacggac gactcccaga tggcgccaa cggtgccaga
1381 atcagaaagc acgtcatgca cctgtttgac tgcgtcgcgg gcgtgctgca cgacactgcac
1441 gtcgtcgtcc tctccttgac ccattccgtac ccaaaacacc tcctcaacga gatgatccac
1501 ggcctcaagc cagtccctt gtgctgctcc ctctcggta tcctgtccac tctgcagaac
1561 ttcgtctgct ctgtccccat ctcgcgggtt agaaagaagc tggaaacgtgc caagcgcaag
1621 ggcattcagcg agtgcaccaat aatcactgca gtaattccctt ttttagcaaca catacttata
1681 tacagcaaca gaccttatgt cttttctctg ctccgatacg ttatccaccc aactttattt
1741 cagtttggc agggggatt tcacaacccg cacgctaaaa attgtattt aactttaaaa
1801 agaacagcca caaataggga actttggctt aaagcaagga ctctccctcc cttatcttga
1861 ccgtgctatt gccatcactg ctacaagact aaatacgtac taatatatgt tttcgtaac
1921 gagaagaaga gctgcgggtg cagctgctgc catggccaca gccacggga cgctgtactg
1981 gatgactagc caaggtgata gcccgttagt gcacaatgac ccgagctaca tggtgcaatt
2041 cccccaccgc gctccaccgc aggtctctag a

//

the above report in format

NCBI Entrez

Nucleotide QUERY

BLAST

Entrez

?

Other Formats: [FASTA](#) [Graphic](#)
 Links: [Protein](#) [Related Sequences](#)

LOCUS PPSURFOP 21839 bp DNA BCT 24-MAY-1994
 DEFINITION *P. pentosaceus* (PPE1.0) sucrose and raffinose operons.
 ACCESSION Z32771
 NID g493728
 KEYWORDS alpha-galactosidase; alpha-glucosidase; enzyme IIabc; fructokinase; insertion element; permease; regulator; sucrase.
 SOURCE *Pediococcus pentosaceus*.
 ORGANISM *Pediococcus pentosaceus*
 Eubacteria; Firmicutes; Low G+C gram-positive bacteria;
 Lactobacillaceae; *Pediococcus*.
 REFERENCE 1 (bases 1 to 21839)
 AUTHORS Leenhouts, K.K.J., Bolhuis, A.A., Kok, J.J. and Venema, G.G.
 TITLE The sucrose and raffinose operons of *Pediococcus pentosaceus* PPE1.0
 JOURNAL Unpublished
 REFERENCE 2 (bases 1 to 21839)
 AUTHORS Leenhouts, K.K.
 TITLE Direct Submission
 JOURNAL Submitted (27-APR-1994) Leenhouts K. K., Institute for Biological Sciences, Department of Genetics, Kerklaan 30, Haren, The Netherlands, 9751 NN
 COMMENT On May 26, 1994 this sequence version replaced gi:[475962](#).
 FEATURES
 source Location/Qualifiers
 1..21839
 /organism="Pediococcus pentosaceus"
 /strain="PPE1.0"
 /db_xref="taxon:1255"
 repeat_unit 487..510
 /note="DR2"
 /citation=[1]
 /rpt_type=DIRECT
 repeat_unit 488..561
 /note="IR1"
 /citation=[1]
 /rpt_type=INVERTED
 repeat_unit 513..561
 /note="DR1"
 /citation=[1]
 /rpt_type=DIRECT
 terminator 2724..2760
 /standard_name="terminator 4"
 /citation=[1]
 gene complement(2929..3864)
 /gene="rafR"
 CDS complement(2929..3762)
 /gene="rafR"
 /standard_name="regulator"
 /function="regulation raffinose-operon"
 /citation=[1]
 /codon_start=1
 /transl_table=11
 /product="regulator"
 /db_xref="PID:g475963"
 /db_xref="SWISS-PROT:P43465"
 /translation="MNGEYKTLANKSFESNVLLFFGQEACLPNYTYKGNVRDSYVIHY
 IQEGKGTFAANHPATVLKAGDIFILPKGTPCFYQADNDQPWKYFWIGFSAGIRIEAM
 LSGSLLAQKCYLRQVQNGHIYADLSELYKVLHIPNSLINDVLLGSLIYRLFYDLLRWY
 PADATNIKVKSTEQFNLAVSYLQENYSTGCTIMDLCHYLNLSRSYLYTLFKTHANTSP
 QKLLTKLRLEDAKQLSTSNNSVQSIANMVGYKDSFTFSKAFKRYSGASPSYYRKSIG

```

I"
-10_signal complement(3837..3842)
/gene="rafR"
/citation=[1]
-35_signal complement(3859..3864)
/gene="rafR"
/citation=[1]
-35_signal 3909..3914
/gene="rafP"
/citation=[1]
gene 3909..5918
/gene="rafP"
-10_signal 3932..3937
/gene="rafP"
/citation=[1]
CDS 3993..5918
/gene="rafP"
/standard_name="permease"
/function="raffinose transport protein"
/citation=[1]
/codon_start=1
/transl_table=11
/product="permease"
/db_xref="PID:g475964"
/db_xref="SWISS-PROT:P43466"
/translation="MQEEHNYKWKVGGRLIYGFAGKNDAFYSILSGYLIIFITSHLFDTGNKALDNRMVSLVTLLIMVLRIELFIDPFIGNAIDRTKNSPGHFRPWVVVGTVSSIILLLFTNLGGLYAKNAMIYLVVFAILYITMDIFYSFKDVGFWMSMLPSLTTDSREREKTATFARLGSTIGGGLVGVLVMPAVIFFSAKATSTGDNRGWFIFALIICLIALISAWGVGLGTREVDSDIRKNKQDTVGVMEIFKALAKNDQLLWAALAYLFYGVGINILGSLEVYFTYIMGKPKFSIILSIINIIFLGLIATSLFPVLSKKFSRKGVFAGCLVFMLGGIAIFTIAGSNLWLVLAAATMFGFPQQMVFVLMVITDSVEYQQLKLGHRDESLALSVRPLIDKFGGAISNGVVGQIAISGMMTGATASSITAAGQLHFKLTMFAFPALMLLIAIGIFSKQIFLTEEKHAEIVAELETRWRTKFDNTTDQVAEKVVTSDLATPIAGQVIPLAQVNDPTFAAGTLDGFAIKPSDGRILAPFDATVRQVFTRHAVGLVGDNGIVLLIHIGLGTVKLRGTGFISYVEEGQHVQQGDELLEFWDPTIKQAGLDDTVIMTVTNSTEFTMMMDWLVKPQAVKATDNILQLHTKA"
gene 5976..8177
/gene="agaR"
CDS 5976..8177
/gene="agaR"
/standard_name="alpha-galactosidase"
/citation=[1]
/codon_start=1
/transl_table=11
/evidence=experimental
/product="alpha-galactosidase"
/db_xref="PID:g475965"
/db_xref="SWISS-PROT:P43467"
/translation="MSLITVDQANRVFHLLNQTLTSYIFAVEQGGTLSHLYFGGHVDHYHGELRYPRVDRGFSGNLPGSTDRTFSRDTLPKEYSTAGEMDYHLPAAIVRHTDGANAYLTYQGYRIEAGKPKLGLPAAFVEDETEAETLTIVLVDQVSQVEFDLQTYTIYRDRPVVTRSVQVCNQGDHAVNLEKVASMQIDFTDQFETITLPGAHANERHPERGSINYGIQTFGSLRGTSQHMNPFLALVDHTTTEFSGDAYGFNLVYSGNHAFELEKDQLDQLHLMVGINSYFNWQLKAGATFQTPEVLMVYTNKGLNAMSQAYHHLIRERVVRSEFKNQERPIVVNNWEATFFDFNEAKLKPIVDEAKQLGIEMFVLDGWFGRDDDNSSLGDWQVDHRKFQGLNHFVKYVHEQGLKFGIWLPEMISYDSKLYQQHPDYLQVPGRSPSPSRNQYILDLGRQAVRNNIFDQLDQLLKSQIDYIKWDMNRHLSDIYSVALPPERQGEVYHRYVLGLYELLERLTTAYPHILFEGCSGGGGRFDAGMAYYMPQIWAISDNTDAVARLTIQYGTSLAYPISLATAHVSVSPNQQTGRETSMSTRSAVAASGVLGYELDLTQLSSADKQIVQKQVQYKQIRPLIQFGEFYRLKSPITSNQAAWMFVSPQQDEAIVMVFNLTSYAQPSLTKTKLVGLNPKLNQNIATKAIFFGDELMQLGFYDPVVYQDYTTKVYHFKAVTEN"
terminator 8265..8289
/standard_name="terminator 3"
/citation=[1]

```

```
repeat_unit 9403..9426
/note="DR2"
/citation=[1]
/rpt_type=DIRECT
gene complement(9519..10424)
/gene="iso-IS30"
CDS complement(9519..10424)
/partial
/gene="iso-IS30"
/standard_name="IS-element"
/function="IS-30 (E.coli) like IS element"
/citation=[1]
/codon_start=1
/transl_table=11
/product="insertion element"
/db_xref="PID:g475966"
/translation="SDAVRTKRSASLKWLPOTHLLGADLSSITYSERIKIETFCEL
GLSNIQMGVRLNRSPSTISYELSRCQPYQAE LAQTD EYKRSQCGRKTKLSDELKQKI
LNHLRLSWSPGMIAHEFKLATKSIYNWLNQGRIGFSLNDLPEHGVQRQRRNVDQRSKYN
QLGRSIEQRPMMFNQRNRIGDFELDTVVGPRGHSKAVLLTLIDRKSRFLWAYRLKDR
TTATVNEALTKFLTTFNGPVHSFTVDRGTEFSGLVSLESQYGIKTYCHAYTPADV
MNALIGIYVVFILKGLVLSTLVLKI"
repeat_unit 10396..10442
/note="DR1"
/citation=[1]
/rpt_type=DIRECT
repeat_unit 10397..10443
/note="IR2"
/citation=[1]
/rpt_type=INVERTED
terminator 10681..10716
/standard_name="terminator 2"
/citation=[1]
gene complement(10727..11593)
/gene="scrK"
CDS complement(10727..11593)
/gene="scrK"
/standard_name="fructokinase"
/function="fructose 6-phosphotransferase"
/citation=[1]
/codon_start=1
/transl_table=11
/product="fructokinase"
/db_xref="PID:g475967"
/db_xref="SWISS-PROT:P43468"
/translation="MLLGAI EAGGTFV CATGAENGQVSDRISIPTTPVETMTAVDD
YFTTHPVD AIGIGSFGPIGVNP HDPKYGYITTPKPGWGD FDFLGH LKSQFNIPLYWT
TDVNEAAYGESMIGIAKDVPNSIYMTIGTGVGAGVISQNHIFNGRTHTELGHMRLNRL
PGDDFKSNC P YHDICLEGIAAGPAVGKRTGKAGKDIPVDDPVWPIITDYIAQACVNLT
VAFAPDKIIILNGGMNQRQLFPMIREKFAAYLNGYEEVPLDDYIVPAGLGNNSGIAG
GLLLAQAAALKNA"
gene complement(13852..15989)
/gene="scrA"
CDS complement(13852..15807)
/gene="scrA"
/standard_name="enzyme IIscr"
/function="enzyme II of the PTS system, sucrose specific"
/citation=[1]
/codon_start=1
/transl_table=11
/evidence=experimental
/product="enzyme IIabc"
/db_xref="PID:g475968"
/db_xref="SWISS-PROT:P43470"
/translation="M NHQEVA DRVLNAIGKNNIQAAAHCATRLRLVIKDESKIDQQAL
```

DDDAADVKGTFETNGQYQIIIGPGDVDKVYDALIVKTGLKEVTPDDIKAVAAAGQNKNP
 LMDFLKVLSDLIFIPIVPALVAGGLLMLANNVLTAEHLFMAKSVEVYPLKGIAEMIN
 AMASAPFTFLPILLGFSATKRGFGNPYLGATMGMIMVPLPSLVNGYSVATTMAAGKMVY
 WNVFGLHVAQAGYQGQVLPLVGVAFILATLEKFFHKHIGAFDFTFTPMFAIVITGFL
 TFTIVGPVLRTVSDALTNGLGLYNSTGWIGMGI FGLLYSAIVITGLHQTFPALIETQL
 LANVAKTGGSFIFPVASMANIGQGAATLAIFFATKSQKQKALTSSAGVSALLGITEPA
 I FGVNLKMKFPFVFAAIASGIAASFLGLFHVLVAMGPASVIGFISIASKSIPAFMLS
 AVISFVVAFIPTFIYAKRTLGDDRDQVKSPAPTSTVINVNDEIISAPVTGASESLKQV
 NDQVFSAEIMKGAAIVPSSDQVVA PADGVITVYDHHAYGIKTTAGAEILIHGLD
 TVNLNGEHFTTNQKGDTVHQGDLLGTFDIAALKAAANYDPTVMLVTNTANYANVERL
 KVTNVQAGEQLVALTAPAAASSVAATTV"
 -10_signal complement(15872..15877)
 /gene="scrA"
 /citation=[1]
 -35_signal complement(15894..15989)
 /gene="scrA"
 /citation=[1]
 gene 15993..17568
 /gene="scrB"
 -35_signal 15993..15998
 /gene="scrB"
 /citation=[1]
 -10_signal 16017..16022
 /gene="scrB"
 /citation=[1]
 CDS 16063..17568
 /gene="scrB"
 /standard_name="sucrose-6-phosphate hydrolase"
 /citation=[1]
 /codon_start=1
 /transl_table=11
 /evidence=experimental
 /product="sucrase"
 /db_xref="PID:g475969"
 /db_xref="SWISS-PROT:P43471"
 /translation="MIWRNKTRYTPYEQWPATKLPQLVAQARQSKWRMQHHIQPTSGL
 LNDPNGFSYFDGQWHLFYQVFPGPVHGLKSWQHVTSKNLVDWHDEGLAIRPDTQYDS
 HGAYTGTALPIDQQLFIMYTGNVRTADWQRESYQLGAWMDTDNHKKLSRPLIAHAPA
 GYTSSFRDPDLIRNDHGYYALIGAQTTTEIGAI LVYFSKDLTTWTCQGELNV PANARG
 YMIECPKSGLDRSTARLIVLSQGLSQATIPYQNIYPNMYLVADQLNIAQAOFTEPHAL
 TQLDDGFDVYATQAINAPDGRALAVSWIGLPEISYPTDRENWAHCLSLVKELTLDGH
 LYQNPVAAVDDLRTTAHDLVFEQQRATVAALNGSFELLTVPADKTVTVNIADQQESG
 QLQVTVDAHGQVMIDRRHTGNSFAEDYQGQTRQVELTAHKTIKIRLIIDVSVFECYID
 NGYSVMTGRFFLNATPSRLNVQGDTTAVTGKVWEWRQSEHTGVDNNETKIK"
 gene 17549..18529
 /gene="scrR"
 CDS 17549..18529
 /gene="scrR"
 /standard_name="regulator"
 /function="regulator of the sucrose operon"
 /citation=[1]
 /codon_start=1
 /transl_table=11
 /product="regulator"
 /db_xref="PID:g475970"
 /db_xref="SWISS-PROT:P43472"
 /translation="MKPKLNDVAKLAGVSATTSRVINNHGYLSSQTKEKVFAAMREL
 HYQPNNMARSLQGKNTRLIGVIFSDISNPFFGELVSRIEKILFAKNYKVILCNSADDP
 QKERDYLQMLMANQVDGI IAGAHNLGIEEYQQYGLPIISFDRLSDNIPIVSSDNYQG
 GWLATQTLHQAGATNVAIFTGKSHAGSPTNGRREGYEAYLTAQQLTPHHELPFELTP
 ALKMMEIKTINTQHQYDGFCSDDLAALLVNVAAQQLSLTVPEQLRVVGYDGTALIRD
 YHSELTTEQPLADISTLILVSLLLQRIEDANCTLESKYTLPVKLIKGFTA"
 gene 18550..20223
 /gene="agl"
 CDS 18550..20223

```

/gene="agl"
/standard_name="alpha-glucosidase"
/citation=[1]
/codon_start=1
/transl_table=11
/evidence=experimental
/product="alpha-glucosidase"
/db_xref="PID:g493729"
/db_xref="SWISS-PROT:P43473"
/translation="MAATIKWWQQAVVYQVYPRSFQDTNHDGIGDLKGITAHDLYLKQ
LGIDVIWLNPPIYRSPNDDNGYDISDYQQIAADFGTMADFDLQAAHDRGLKIIIMDLV
VNHTSDEHPRFKRSRQDRTNQYRDFYFWRSGNGKKAPNNWEAAFGGSAWQYDEQTOQQY
YLHTFSTKQPDLNWENPTLRESVTMMTWWLNKGVDGFRMDVINQISKLPGLPDCPLK
PHSQFGDARVTNGPRVHEFLQEMNQEVLSQLFDIMTVGETHGVTPADALKYAGADQHEL
DMVFEFQHRLDQSQHGLGWSTRKTPVALKKVISDWQVGLEGRAWNSLFWNNHDTP
RAVSRFGDDRPAYRVRSAKMLATCLHLLQGTPYIYQGEELGMTDAHFTELASYRDIES
LSAYRDLVTERQLLSPADMMARLAAASRDN RTPMQWDTEVNAGFSDAAPWLTVNPNY
RQINAALADPDSVWYYYQHLIQLRQYPSVTLSFELLWADDPQYSYMHGNGKADL
ASLLQFHRSRDTVPTTGSISDPTAKCLISNYGEQQPNKLRPYEAWVYQLA"
terminator 20341..20389
/citation=[1]
gene 21090..21677
/gene="iso-IS3"
CDS 21090..21677
/gene="iso-IS3"
/standard_name="IS element"
/function="IS3 (E.coli) and IS981 (L.lactis) like IS
element"
/citation=[1]
/codon_start=1
/transl_table=11
/product="insertion element"
/db_xref="PID:g475971"
/translation="MCRILGVSRQAQYYRYRSPKPSKRRAEDADLKQRLRIFAEFKQR
YGVMKIHHELNLELQPLQLRCSPRRISRLMKELDIHSVTVNWKAAASASKTKVEQRPN
LLKQDFSTTGLNQKWTADMTYIQTKRNGWCYLSTIMDLHSRRIIGYSFSKKMATDLVL
KTLESAVKNRTITGGPDYPYGFRTGYQRLITINV"
BASE COUNT      6181 a      4374 c      5015 g      6269 t
ORIGIN
  1 gcatgcttga  ctgggtgctt agccgcaaat gatcaggaga  gaagcatcg  gactaggaaa
  61 ggagggcaac acggttgctt taatcaacaa agaccaggc  gatagttca  tcttcctgtt
121 ttagttgac  agataaatga  cgattaaagg  cggccaccag  tttgcgtgcg  aaggccatct
181 ttcgtcgaa  ttccgtcatg  tgcgaagtaa acgggacgct  ttggtaatta  aagttttcgc
241 cgatatagcc  cggatcagtc  acgtacgc  agttgtttc  gtcgtggcg  ctggcccggtt
301 tcaggacggt  atccaaatac  tgatagagaa  tcgttggatc  gtagttcagc  ggcttatcaa
361 gggtcgtgga  aagctgttag  ccgggtgctt gaccagcgaa  ttgttcgtt  atttagactt
421 gggtaaggct  ttgatacaat ttcaaaaacga  tcactcctct  tcttttcaact  ttttagtata
481 cagttatgg  agattgtaaa  attaatccga  acgctgatct  tttttgtccg  aacagcggtt
541 ggattaaattt tacaatctac  cttttaccaa  gcaattgtatc  aaatgctac  taacgctagt
601 cctagtgtata  aattgtctaa acgcttgc  gcaatcttt  cccaaagcatc  tatttctcg
661 cgtgatatgt  tgggtgcaat gattggcg  gtatgaatat  tgctacagct  aaaagtgaat
721 catttgcagt  taatgaattc  aaacagcagt  atatcgatgg  ctataataat  ttaagcccta
781 ataaactatc  aacagttcca  gataaaagg  ccaaccaac  tgagtattct  caacgaattt
841 gggtgcattgg  cgacgtcatg  acgacacgaa  tggaaagaaac  cctgaacaag  ggtttgagtc
901 gcccgcattaa  aaaatcagac  atcaataaaa  tgaccggc  tattcctcaa  agcggtgatc
961 gaatttgatga  caatttagca  acgcctatga  accaaatgct  atccagaatt  catabgttac
1021 caactaatga  agctatacgc  aatagcaac  gtggtaagaa  tagaaacgtat  gatgaacaga
1081 atgttcagta  tgtgtatgtt  ttaaccgagg  aagattgtatc  gtgtttgtat  tatttgtgaa
1141 ccacttgata  gacaaacatt  tgcctatgg  caagcaccga  ttccagtcac  ggatactcat
1201 ccaagggtgtc  gttggccagct  ggttagcctgt  gatgaagatg  ggaatctgt  gtcgtggtaa
1261 cttgtatagg  tatttaagga  acaaaaaaaag  cccacttagt  taaaactaa  gtgggtcagg
1321 aatattatta  attggcttcg  ctcacaacaa  agtcgagg  tgctaactaa  gttagcatct
1381 ataatttagt  ataagaacct  taaatgaac  ttaaaaacta  attctttgtat  ttaccgacaa
1441 agaaagaaac  tacagcaaca  acaattactg  ctccaataat  tgaaggata  attgcccattc
1501 cggctaattt  agggccccaa  gatccgagta  tagcttctcc  aatagaggaa  ccaactaattc

```

1561 cgccaacaat attagcaatc caccatttgc atttccctt gcttagtaatt gctccagcaa
1621 tagcaccaat gatggacca ataattaaaa cccataacca atgcataaag catcatcctc
1681 tcttattttt cgagggatgt ttttagatt tgttatagtt ttgaatttct ttgaaaatgg
1741 ctaataaaaag tttagatata gaaatgataa ttgataatt tgaccaaaatg ccaaatctt
1801 tcataatgaaa tcctccatta tctaaattgc attcttctaa acctaattt aggcaattga
1861 atatttgcatac acaaggattt cgattgtgcc attcttttta aaataggaat ttttatttt
1921 ccttttctc aattacagac gttaaagaat aattgattt gttattgcac ttacaaacaa
1981 ttttagtctac ggactaaaa aggagttt gatgaaaa caaatttgca attttttgc
2041 gatcctaacc cagaacctaa gccaaacgat tctaaccctt atcctgatgt gacctttgg
2101 gctgtgtgt atagacggtt cttgcataa gacggaaact tcttcattt tttgtgttag
2161 gaatttcaat ggtacctgaa ttttgcattt tggtgtctt ttttaccat ttttagtaca
2221 tggcttaactt aattataaaa cgcgcgagaa taaatttttgc gggaaaaag gaattaaaaa
2281 ccataataaa ttttattctat tgaatttaaa tgactatcg tttgtcaatg taagatatcc
2341 agtaactcaa gcttagttga atgaaatgtt aatcaaactt ggtatgaag ccgatattac
2401 tgcgcctaaa gagatgcagt ggtcgatgtt tcatttgcgg catacagttt caacaagact
2461 tggaaaata cagccggat gagttatccg tggcgccag cgcgtatggg acatacgta
2521 gaagagttca tgcagactta ttttgcattt gatcagaatc gttcgatgtt aatgtgtgc
2581 aagtgggtat gattgactaa atttctcaaa cacacacacg aaacacacag aactcaaaga
2641 taattaaata aacgcgattt ctgcgggatt tcgaagctca actttgtatc ttgcagacgc
2701 ataaacaaaaa taattttat ttgcacaaagc cgacaatagc gctgttatttgc tcggtttttgc
2761 ttgtttaaa aaaggctgta tcaatttttgc tcaacacaca gcaaaatttgc aaaaagtcgc
2821 gcgttatttgc ggtgttccca taatttttgc aggacaactt ttcatttgcattt aattcgatag
2881 cattagcaga aagacaaatc acacgggcaaa catagtggc attatgattt agattccat
2941 gcttttacga taatagctgg gcttagcacc acttgcgtt ttttttttttgc tggagaaatg
3001 gaagctatcc ttgttagccta ccatatttgc aatgttttgc actgaattt tactgtgt
3061 tagacgctgt tttagcatctt ctaagcgcattt ttttagtttgc aatgttttgc
3121 agcgtgcgtt taaaacaagg ttttttttttgc gtttgcattt aatatttttttgc
3181 gtccataatg gtacaaccatg tactataatt ttcttgcatttgc taagaaacag ctaatttttgc
3241 ttgttcagttt gacttactt ttatgttgc ggcgttgc ggttgcatttgc gcaataatgc
3301 atagaacaat cgataagatca aagaccctaa tagaacatca ttttgcatttgc aatttttttgc
3361 gtgtaaaactt ttataaaggat cagagatgc ggcgttgcatttgc ttttttttttgc
3421 taatgtatc ttcttgcatttgc ataaaggatc agaaacgc gtttgcatttgc ggataccggc
3481 gggaaaaaccg atccaaaatgtt ttttttttttgc ttgttgcatttgc ttttttttttgc
3541 ttgttttttttgc ggttgcatttgc aatattatgc ggcgttgcatttgc gatgggttgc
3601 tgcagcaatg gtacccatttgc ctttgcatttgc gtttgcatttgc acatagtttgc
3661 attacccatttgc taatgttgcatttgc ttgttgcatttgc agcccttgc
3721 ttgttttttttgc ttatgcatttgc aatatttttttgc ttgttgcatttgc
3781 taaaatatttttttgc ttgttgcatttgc aatatttttttgc attaatttttttgc
3841 tactgttttttttgc ttttttttttgc tactaatttttttgc aataacaaaaa aatttttttttgc
3901 gattactatg ggcacccaaatg tagacaggaa ttataatgc
3961 acaaaaaatgtt ttgttttttttgc ggttgcatttgc ttatgcatttgc
4021 taggaggttgc ctttgcatttgc gtttgcatttgc
4081 ttttttttttgc ttatgcatttgc
4141 tagataatgc aatgttgcatttgc ttgttgcatttgc
4201 ttttttttttgc ttatgcatttgc
4261 ttttttttttgc ttatgcatttgc
4321 ctttgcatttgc ttatgcatttgc
4381 ttttttttttgc ttatgcatttgc
4441 ttttttttttgc ttatgcatttgc
4501 ctttgcatttgc ttatgcatttgc
4561 ctttgcatttgc ttatgcatttgc
4621 ttttttttttgc ttatgcatttgc
4681 ttttttttttgc ttatgcatttgc
4741 atgttgcatttgc ttatgcatttgc
4801 taggttgcatttgc ttatgcatttgc
4861 ttgttttttttgc ttatgcatttgc
4921 ctttgcatttgc ttatgcatttgc
4981 ttttttttttgc ttatgcatttgc
5041 ttttttttttgc ttatgcatttgc
5101 aatgttttttttgc ttatgcatttgc
5161 ttttttttttgc ttatgcatttgc
5221 ttttttttttgc ttatgcatttgc
5281 aatgttttttttgc ttatgcatttgc
5341 agcaaatatttttttgc ttatgcatttgc
5401 ggagaacaaaaa gtttgcatttgc ttatgcatttgc

5461 atttagctac accaatcgct gggcaagtga ttccacttgc ccaagtcaat gatccaactt
5521 ttgcggctgg aacgttaggt gacggatttg ctattaaacc tagtgatggc cgaatattag
5581 ctccatttga tgcaacggta cgtcaagtat ttaccacacg acatgcagtt ggcttagtcg
5641 gtgataatgg gatcgctta ttgatccata ttgggttggg aactgttaaa ctttagggaa
5701 cgggatttat ttcttatgtt gaggaggggc agcatgtaca acaaggggat gaattacttg
5761 agtttggga tccaacgatc aaacaagctg gtttagatga tacggttatt atgacagtga
5821 ctaattcaac cgaattcaact atgatggatt gtttagtcaa gccaggtcaa gccgttaaag
5881 caactgataa tattttacag ttgcatacta aagcataatg tgccacttgtt tatcatttaa
5941 tttaaaatt agtaattccc cgaggaggaa ataaaatgtc attaattacg gttgatcaag
6001 cgaatcgctg tttcatttgc cataatcaaa ctcttctta tatcttgca gttgagcagg
6061 gcggtacttt aagtcatatcattttgggtt ggcatgtgga ccattaccac ggtgaattgc
6121 gctatccacg agtcgatcggtt gtttttccg gtaatttacc gggatcgact gatcgactt
6181 tttcacgtga tactttgccc aaagaataca gtacggctgg taaaatggat tatcacttac
6241 cagcagcaat cggtcgcat actgatgggg ctaatgcatt atacttagtt tatcagggat
6301 atcggatcga agcggtaaa ccaaagttaa gcgggctacc ggcagcttt gtggaggatg
6361 aaacagaggc agaaaactttg actatcgat tagtgatca agtaagccaa gtcgagttt
6421 atctacaata tacgatctac cgtgatcgac cagttgtcac ggcgtcgggtaaagtttacgg
6481 atcaaggtga tcatgcagtt aatttagaaa aagtgtctt aatgcaaaatc gattttacgg
6541 atcgtagtt cggaaacaattt acattacctg gagcgcatgc caatgaacgc catccagagc
6601 ggggttcgat caattatggaa atacagactt ttggtagttt gcccgtact tccagtcatt
6661 aaatgaatcc attttggcc ttgggtgatc acactacgac agagtttagc ggtgatgcat
6721 atggttttaa cttggcttat tctggcaatc acgcatttga actggaaaag gatcaacttg
6781 accaacttca tctaattggc gtttataattt caattggcaaa cttaaagctg
6841 ggcgcacttt cggaaacacca gaagtattaa tggcttatac aaataaaggt ttgaacgcga
6901 tgagccaagc ctaccaccaat ttgatccgtt aacgagtagt acgttagtgat ttttttttt
6961 aagagcgtcc gatttagttt aacaactggg aagcaacattt ctttgat ttttttttt
7021 aattgaagcc gatcggtt gaaagccaaacg agtttaggcatttttttttttttttttttttt
7081 atgggttgggtt tggacatcgatgatgatatttttttttttttttttttttttttttttttttt
7141 atagaaaattt tccgcaggcc ttaatcatttttttttttttttttttttttttttttttttttt
7201 aattttggat ttt
7261 atccagatatt
7321 ttttagatctt tggctggcaaa ggcgtacgaa ataatatttt tgaccagctt gatcaactt
7381 taaaatcgaa gcaaaattgtat ttt
7441 atcccggttgc ttttaccacca gaacgcggggttgcgttgcgttgcgttgcgttgcgttgc
7501 ttt
7561 caggtgggttgcgttgcgttgcgttgcgttgcgttgcgttgcgttgcgttgcgttgcgttgc
7621 gtgataacac ttt
7681 caatttcgtt agtaccggcc catgtctcgttgcgttgcgttgcgttgcgttgcgttgc
7741 cgtcaatgtc gactagaagt gcaatggcc gcaatgggttgcgttgcgttgcgttgcgttgc
7801 taacacaactt ctt
7861 agatttagacc actgatttcaat ttgggtgat ttttttttttttttttttttttttttttt
7921 atcaagcggc atggatgtttt ttgggttgcgttgcgttgcgttgcgttgcgttgcgttgc
7981 atctgacatc ttt
8041 agctgaatt
8101 ttggcttcttt
8161 ctt
8221 gtacacagac gaaaggggctc atgggagtttttttttttttttttttttttttttttttt
8281 gcaatggccatc ttt
8341 gggagatgggttgcgttgcgttgcgttgcgttgcgttgcgttgcgttgcgttgcgttgc
8401 gatt
8461 ctt
8521 ttt
8581 ttt
8641 acactctcgatc ttt
8701 ttt
8761 ttt
8821 atcttccatc ttt
8881 ggg
8941 ttt
9001 ctt
9061 ttt
9121 ttt
9181 ctt
9241 aggg
9301 aggg

9361 gtggctaaact taattataaa acgcgcggat aattttgata attggtagat tgcaaaattta
9421 atccgaattt ttggacaaat ttgtcagcat aacctgatac ggtgtttgct agtcaggtat
9481 ttaagcggt cgctggttaa ttggagtaa cgtcgtcgtt aatctttag tactaatgtg
9541 ctcaaaaacga gtcctttag gataaaaaca acgtaaattc cgattaaagc gttcattact
9601 accacgtca gtcggagtata agcatggcag taataggctc taataccata ttgtgattca
9661 agtgataacta gaccgctaaa ctcagtgcctt cggccacccg tgaagctgtg caccggacca
9721 ttaaaagtgg ttagaactt agttagggct tcattaaacag tcgctgtcgt ccgtcttt
9781 aaccggtagg cccaaaggaa cctgtatct agtcaaaaat aactgcctt
9841 ctatccccac gaggaccaac ttcaattgtat cgcggccaaag cggcgatgcg attacgttg
9901 ttaaacatca tggacgctg gctgtacgcca tggtcaggta gatcattcaa ggagaaacca
9961 tggtaacgt tacggcttg attataataa gatttagtag ctgtttaaa ttctgtgagca
10021 attctccctt gatttagcca tagacgtaaa tggttgagaa atttttgctt taactcatcg
10081 atcatccctg gtgaccagct acattgtgtat cgctgttattt cggcatctgt ttgtgtaat
10141 ctcagcttag tttccgacc tcagcctgtat aagggtgaca tcgagataat ttttgacgg tgatcggttc
10201 tcagcctgtat aagggtgaca attggacagc cctagttcac attttttttttt gatcattcaa ggagaaacca
10261 agccgaacgc ccattggat 10321 cgttccggat aggttataact agacaaaaga tcagctctt
10381 caccattttt aggaagctga ttttttgc cgaacagcgt
10441 ccatgtacgt gtcgtatatt cgcggaaatca tttcgaattt
10501 cgctacaacc actctggca atgattaatg ggatattatg
10561 agacttataa ggattggccc gttattattt caaatgttcc
10621 tcaactgtttt aaccgtatata atgcattaaa cgaatgtac
10681 ccgctaatac gtttaaacac gtatcggcgg actttttgc
10741 aagcagccgt agctaacaat aggccacccg caatgcaga
10801 gaacgatata gtcatccagt ggtgggactt cttcgttagc
10861 tttcacgaat catggaaag agtgcggcgtt gattcataac
10921 ttcagggtgc gaaggcgaac gtgaggttga ccatgcctg
10981 ttggccagac cggatcgta acggaaatat cttcccgac
11041 ctgctggccc agcagctaaa cttctaaac agatgtcatg
11101 aatcgcccc aggttaaccgg ttgagtcgc tggtaccaag
11161 taaaaatgtg gttctgacta atgacggccag cgcacacc
11221 tgctgttagg gacgtcctt gcgattccaa tcattgattc
11281 catcagtgtt ccagtagat gggatgtga attgacttt
11341 agtcacccca gcccggtttgg gtagtcgtgg taatgtaa
11401 taacggcgtat ggggccaaat gaaccgatac caatcgctc
11461 agtcgtctac cgcagtcgt gttcaacccg ggggtgtcgt
11521 taacttgccc gtttcagcg cccgtggcac aaacaaattt
11581 cacctaaaag catagtgtgc ctcctgtatg taaaatgt
11641 attcttaattt cgaaattttt ttaacgatta cattaatttt
11701 ctatgtaaat cagcgttggt ggtttaagt ggaatcccc
11761 gtatagcgat taccgatga atcgctatac gctaattcag
11821 aatgggtgtct gagtttgc acgagaatta agcccat
11881 ttttatctg ccgttactat caaccacgc acaacgtgtc
11941 aagtggccct tgataaaaatc agatttcaatc
12001 agttgtgtt cactaggac gtttgcgtt ggtttaatc
12061 caggctaatt gtaatcgat tcaaatggt gtattcggc
12121 gcggttatata ggtgtctgat tccgtggt ggtttaatc
12181 gttgctcggt cgactggatc tccgtggt ggtttaatc
12241 ataccaaaaat caaaccggcc accaccagct
12301 gcgcgttca agcgtgttag gatatcataa agtccac
12361 ccctgttgcg cagccggtaa gatcgatcg
12421 catttaaat aatcgagttt gatcgatcg
12481 tcacggacag cggctgcga ccatcgat
12541 cgatcaacat attgtataac gatcgatcg
12601 atcatctcg gttccacccaa gagaccaat
12661 agggggtca gaccattagg ttttgcgtt ggtttaatc
12721 ttttcgcgt tacgggtgac tccgtggt ggtttaatc
12781 tgggggggt gttcgattaa tccgtggt ggtttaatc
12841 tcccaagttt taagcacact gatcgatcg
12901 tcgtgaaata cttcgacat gatcgatcg
12961 ggagtctgaa aactgttattt gatcgatcg
13021 attgttaggc ggctatgtt gatcgatcg
13081 catgctagtg cacaaccaat aacagtccca
13141 gccatgaacg gttgggttgc gatcgatcg
13201 cctgggtqta aacggtaaa gatcgatcg

13261 agcgcagttt attggcgtt tggtagatct agttgagcgc tggatagggc tgtaacttgc
13321 aaatttagtag taccgtgatg gcgccaggtt gtgcctcgca agataagccg ctgatTTCA
13381 aaaatagtgt agttaagatc catctgttac ttggctactg catcggttca atgaatgact
13441 aggggttta caggcgtgtt tttgcttacg gtgggtggca attggcgcga gttgattggc
13501 tgcatttca cagaaaaacc agtatattct aaaattggca gtaatttgatt gttagcatcc
13561 ttaataacgt aagctggctg gcgatgtca ccactgccaa tagtggaaa ttccgagtggc
13621 agtgaagtca cggaaatatgg aaagcgttct gttgggtcat tagcaaaagc atggctaccg
13681 gatggcatcg gctcaaaatg cgggttgggg gaaaagtaac gtccaaaata tcgatgcact
13741 gggtaacgtt gagctaaaat ttgaataaca tagctaattt ggtcattatg aagattgatt
13801 aagccggttt tggggtcata ttcaacaggc atgacgttct cctaataaaag ttttagacagt
13861 ttagcagca actgagctag ctgcgggagc agtcaacgc acgagctgtt caccagcttgc
13921 gacattcgtt actttgagtc gttcaacattt agcgttattt gccgttattt tgacgatttag
13981 catgacggtt ggtatcatgt tagcggctt taaggcagcg atgtcaaaacg ttccgagtag
14041 atcaccttga tgaacggat cgccttctg aacgttagt gtaagtgtt caccattaa
14101 attgaccgtt tctaattccta agtgaatcag aatttccggca ccagcgtt ttttgatgcc
14161 ataagcatga tggctatcat aagtgcacgt gatgaccccg tcagctggc cgacgaccctg
14221 atcagagctt ggcacgattt cggcgcctt ccccatgatt tcggctgaga aaacttggc
14281 gtaacttgc ttgagacttt cgcttagcccc cgttaccggc gcaactgataa tctcatcatt
14341 aacattgtatc acggtaactcg ttgggtctgg tgattttact tgatcacgt cgtcaccggag
14401 tggctgttgc gcatagataa aagtggaaat aaatgcgacg acgaacgaga tgacagcact
14461 gagcatgaat gctggaaatcg actttgaagc aatcgaaatg aatccgatga ctgatgcggg
14521 acccatcgcc acggacagaa cgtggaaatag tcctaagaaaa gctgaagcaaa ttccctgaggg
14581 aatcgctgca aagacaaaatg gaaatttcat cttgagggtt accccggaaaa tagcagggtt
14641 cgttaattccg agtaacgcgt agacccggc tgaagaatgc agggctttt gctttggct
14701 cttagtgccaa aagaaaaatag ctaaagtggc ggcacccgtc ccaatgttgg ccatcgaggc
14761 gactggaaatg ataaaggaaac caccagttt ggcaacattt gccaacaact gggtttcgat
14821 tgcggggaaa gtttgcgtt gaccgttaat aacaatttgcgaaataa aaccaaaagat
14881 tcccatatcca atccagccgg tgcgttata taaggctact aaaccgttag ttaatgcac
14941 gctcaccgtt cgttaagacgg gaccaacgt tgtaaaaatggg agggaaaccag taatcacaat
15001 ggcacacatc ggggttaaaacg tgaagtcaaa tgcccttta atgtgtttt ggaagaattt
15061 ttcaagggtt gcttagatga aggccacggc taagactggg agcacttggc cttgatagcc
15121 ggctgtgca acgtgttacc caaagacgtt ccagtaaaacc atcttgcgg ctgcctatgg
15181 ctttgcacgt ctatacggt taacttagtga tggtaagacc atgatcatgc ccatcgtagc
15241 acccagatag ggggttccac cgaaggcgtt cgttgcgtaa aatcctaaga ggattggcaa
15301 gaaagtaaac ggcgcacttgc ccatcggtt aatcatttgcgacccatc ttggccagg
15361 gtaaaacttca acaactgtact tcgcctataaa aagatgtcg gcaactaaaaa cgttgggttag
15421 tgcctttaat agaccccccgg caacttagtgc agggacgattt ggaataaaaaa tatcagataa
15481 gactttgagg aagtccatta atggattttt attttgcaccc gcaacggccaa ctgccttgc
15541 gtcattcggtt gtcatttctt taagacctgtt tttaacgattt aaggcgttcat agactttatc
15601 gacatcgcca gggcaataa tgatttggta ctggccgtt gtttgcagg tcccttaac
15661 gtcgcgtcg tcatcttgcg cttgttgcattt aatcttgcattt tcatttcgttgc tgaccaaaacg
15721 gaggcgtgtc gacactgggg cggcggctt aatgtgtttt ttaccaattt cattaagtac
15781 gcggtcggca acttcttgcattt gattcataat aatttactcc ctgagttcaaa aattttttt
15841 aatttataaa tgcacgttctt acatggaccc aataatagca attttggaaa atatgtcaag
15901 ctttgcatac aaaattggat tgcgttgcataa tctgtttttt aagccgaattt tggaaacgtcg
15961 acatgaaaat ataaatataaa aatgtcaaaac gatttgcataa tgcgtgttgc tgggtttata
16021 atagtgcacaa ttaatgcgaa ctttgcgtt aatgcgttcatc tcatgtatgcg gaatcgtaaa
16081 acccggttata ccccttatgtt acatggccca gcaactaaggc taccacagct cgttggccagg
16141 gacacggccatg ctaatggcg gatgcacat catattcgc ctacatcagg tttgtttaaat
16201 gaccccaatg gctttcata ctttgcgtt ctttgcgtt ctttgcgtt ctttgcgtt
16261 ttccgtccgg tacacggact gaatgcgttca acacacgtt cctcttgcataa ctttgggttgc
16321 tggcatgtatc agggcttagtgc gatttgcacca gatacggcgtt acgactcaca cggcgcgtac
16381 acgggtacag cactaccaat tgcgttgcattt ttattttttt tgcgttgcggg taatgtccgt
16441 actgcccactt ggcacacgttca atgcgttgcattt tttagtgcctt ggttgcacac ggataatcac
16501 attaaaaatgc tgcgttgcgttca actaatttgcg catgttccgg ctgggttacac gagttcggtt
16561 cttgtatccgg acttgcgttca gatgcgttca ggttgcgttca ggttgcgttca ggttgcgtt
16621 actactgaga tgcgttgcattt ctttgcgttca ttcttgcattt atcttgcacac ctggacgtgt
16681 caaggcgtgtt tgcgttgcattt ctttgcgttca ctttgcgttca tgcgttgcattt
16741 ggtctggatc gatcaacacgc ctttgcgttca ttcttgcattt ggttgcgttca ggttgcgtt
16801 atcccgatc agaatttttgc tccgttgcattt tttttttttt tttttttttt
16861 caggcacaat tcaatggccca ccacggccctt acgttgcattt atcttgcacac ctggacgtgt
16921 gccacgcagg caatcaatgc gccggatggc ctttgcgttca ctttgcgttca tttttttttt
16981 ctttgcgttca tttttttttt tttttttttt tttttttttt
17041 gaactaacgc ttaatggatgg gcacccatgttca aatggggccca attttttttt
17101 cggacgcacgg ctttgcgttca tttttttttt tttttttttt

17161 ggcagcttg aactcttact gacggttccc gctgataaaa cggtcacggtaacattgct
17221 gatcagcagg aaagtggcca gctgcagggt actgttgatg ccaaccatgg tcaggttatg
17281 attgatcgcc ggcatacggg gaattcggtt gctgaagatt atggtcagac tcgtcaagtt
17341 gagttaacgg cccataagac gataaaaatc cggctaatca tagatgtgtc agtcttggaa
17401 tgctatattt ataacggcta ttccgtgatg accggccgtt tcttcttaa cgctaccccg
17461 agtcgtctta acgttcaagg tgacactacg gcagttaccg gtaaagtctg ggaatggcgc
17521 caatcagagc atactggagt cgataacaat gaaaccaaaa ttaaatgatg ttgcgaagct
17581 agcgggggtc tcagcgacaa cgggttctcg gtttattaaat aaccatggct accttagtag
17641 tcagactaaa gaaaaagtt tcgcggcgat gcgagaactg cattatcagc cgaataatat
17701 ggcacgctca ctgcaaggaa agaatacgcg cttattggc gtcattttt cagatattag
17761 taatccgttc tttggtgaac tgggttcccg gattgaaaaa atcttattttt ctaaaaatatt
17821 taaggtgatt ttatgtataa gtgcggatga tccgcaaaaa gaacgcgact acttacaaat
17881 gttgatggct aatcaggtcg acgggattat tgctgggtcc cacaatttag gtattgaaga
17941 atatcaacaa tatggcttc cgattatttc atttgaccgg tacttatccg acaatattcc
18001 catcgtagt tcggataatt accagggggg ctggctagca acccagacat tgcacccggc
18061 gggggcgact aacgtggcta ttttactgg taagtacac gctggttcgc caacgaatgg
18121 tcgccgtgaa ggatatgaag cctacttgcac tgcccaacag ctaacacccc atgtccatga
18181 gtaccgtt gaattgaccc ctgcgctgaa gatgatggaa attaagacga ttatgacaca
18241 acatcagttt gatggatttt tttgtatgtgatgatggatggcc gtttacttag tgctcaacgt
18301 cgcgcacaa ctgtcattaa cgggttctcg gcaattacgc gttgttgggtt atgatggcac
18361 ggcgctcata cgtgattatc attctgaaat aacaacagta gaacaaccgt tagcggatata
18421 tagcacgtt ctggttctc tactatttgcg gcggttggat gatgcgaaatt gtacactgg
18481 atcaaaaatat acgttaccgg ttaagctcat caagggtttt accgcgttaac taaggagtt
18541 atcgtaatta tggcagcaac aatcaagtgg tggcaacaag cgggttctta tcaggtcttat
18601 ccacggagtt ttcaagatac caatcatgat ggttgggtt actttaagggg gatcacggcc
18661 caccttagact acttaaagca attaggaatc gacgtcattt ggctcaatcc catttatcg
18721 tcacccaacg atgataatgg ttatgatatt agtgattatc aacaaatttc ggcggacttt
18781 ggcactatgg ctgatttga tgaactatgg caagggccc atgaccgggg cttaaagatc
18841 atcatggact tgggtttaa ccataccctt gatgagcatc cttaggttcaa acgcagtcgc
18901 caggaccgca ccaatcaata tcgtgatttc tactttggc gctcaggtaa cggcaaaaag
18961 ggcaccaata attggggggc ggctttgtt ggtttagcat ggcaatatga tgagcagacc
19021 cagcagtattt atttacacac gtttcaaca aaacaacctg atttaaactg ggagaatccg
19081 accttacgag agtcgttta tacgtatgtg accttgggtt gtaacaaggg tggggacgg
19141 tttcgaatgg acgtcattaa ccaatcttctt aagtaccgg gattaccaga tggccactg
19201 aaaccgcaca gtcagtttgg cgatgtctcg gtaacgaacg gaccgcgagttt tcatgaattc
19261 ttgcaagaaa tgaaccaaga agtattgtca cagtttgata tcatgacggg tggggagacc
19321 cacgggggtga caccacggc tgcgtgaag tatgcggggc ctgatcagca cgaatttagat
19381 atggcttttgc aatttcaaca tctgcgactt gataatagtc agcatggct tggtaagtgg
19441 agtacgcgtt agacaccgtt agtggcgctt aagaaagtca ttatgtactg gcaagtcgg
19501 ctgaaaggc gggcttggaa cagcctgttt tggacaatc atgataccgc acgagcggc
19561 tcacgttttgc gtatgacccg accagcatat cgggtacgtt cggctaaat gctcgcaacg
19621 ttttacacc tcttacaggc aaccccatat atttaccaag gtgagggact gggatgact
19681 gacgctcatt ttactgaact agccagttac cgagatattt aatcgctcag tgcctatcg
19741 gatgttgcgatc ctgagcgcca gtcgtatc cggcagata tggatggcccg attagcggc
19801 gcttctcggtt acaactcgcg aacggccatg caatgggaca cggaggttaa cgggggttt
19861 agtgcgcgac cgcgcgttcaat caccgtcaat ccaactatc gtcaatcaa tgcgtccgc
19921 gcttggcttgc atccggattt agtgggttac tattatcgc acttgcattca attacgcac
19981 caataccctt cagtgcgtt aggttgcgtt gagctttgtt gggccgtatga tccgcgttat
20041 tcttatatgc acggcaatgg gaaggccggc ctggctatgtt tggatggcccg attacgc
20101 gacactgtcc cgaccactgg atcaatatct gacccacgg ccaaatgtt gattgtat
20161 tatggcgacg agcggccaaa taaattacgg cttatgggg cttgggttca ccaactcgca
20221 taggttattca ggacttggctt aattggggat gtcgttggaaat tatacccaat cgttacgatt
20281 agtgtgaattt ggggttgcgaaat tcttgcgtt atggctttaa taagcgacca cctgactaaa
20341 aaaactgtttt tcaaaaggat aactttttttt aagcgtttttt ttagccggcgg cgaccacttg
20401 gcccacgttgc ggctcatctt gtcgttgggtt accttgcattt aagcccgtt ggttacccgg
20461 taggttgcgttgc tgaccaatgg accgttatctt agcacttgc atagtaattt
20521 taaggccggcgt tgcgttgcgtt aatggatggat gtcgttggat gcttggaaac gaattgtt
20581 actgttggcttgc ggcaagtgcg gggcgatgtt aacggccatggc gcatcaattt tggccattgc
20641 taagtcatctt tggctccaaat cccatcgatca gtcgttgggtt accttgcattt aatcgatgg
20701 gggcttggat gtaagctgtt ttcgttgcgtt aacttttgcgactt ggggttataa tgaataat
20761 gtcgttgcgtt aatggatggat ttttgcgtt aacttgcgtt cggatggat gtcgttggat
20821 acattatcaat cctatataatgc caaggccatggcgtt cggatggat gtcgttggat
20881 tggcttgcgtt aacggccatggcgtt aacttgcgtt cggatggat gtcgttggat
20941 cgccacgttgc aatcaaaggcg atggaaaaggc gacggccgtt gtcgttgcgtt gagaacgg
21001 tcctaaaaaa gcccgttgcgtt tccttgcgtt aatcgatggat ttcgttgcgtt

21061 caagaagcca tcaccaccag gtaaccaaga tgtgccgaat cctcggtgtt tccagagctc
21121 agtattatcg ttatcgatcc cccaaacctt caaaacgccc ggccgaagat gcggacttga
21181 aacaacggat tctgcggatc tttgcggaaat ttaagcagcg atacgggttt atgaagatcc
21241 accatgaatt gaatctggaa cttcaaccac tgcaagttcg gtgcagtcca cgacggattt
21301 cccggctcat gaaggaactg gatatccact ccgttaccgt caataagtgg aaagcggctt
21361 cggcttccaa aaccaagggtt gaacagcgtc ccaacttgct taagcaggtt ttctcgacca
21421 ctggtttaaa tcaaaaatgg accgctgata tgacctatat tcaaacgaag cgtaatggct
21481 ggtgttactt atcaaccatc atggacctgc actcacgacg gattatcgac tattcgttct
21541 caaaaaagat ggctactgat ttagtcttaa agaccctgga aagcgcgggtt aaaaatcgaa
21601 ccattactgg tggacctgat tatccatacg gatttaggat cacaggatac cagcgttattga
21661 ttacaatcaa cgtttaactg agctacatata ccgccactca tacagccgtt aggttatccg
21721 tatgataatg cgccaataga atccttcac gcttccctca aaaaggaatg tgtttatcca
21781 gtgccggctt ttgaagatta tgaaactgtc gctgccgtcc ttttgaata tgtcatgc

//

Save the above report in format

NCBI Entrez Nucleotide QUERY
BLAST Entrez ?
Other Formats: **FASTA** **Graphic**
Links: **Protein** **Related Sequences**

LOCUS PDCRAFOPER 21838 bp DNA BCT 23-MAY-1994
DEFINITION *Pediococcus pentosaceus raffinose operon genes.*
ACCESSION L32093
NID g493181
KEYWORDS agaR gene; agaS gene; agl gene; alpha-galactosidase; alpha-glucosidase; fructokinase; insertion element; permease; rafP gene; rafR gene; raffinose operon; regulatory protein; scrA gene; scrB gene; scrK gene; scrR gene; sucrase; sucrose-6-phosphate; transport protein.
SOURCE *Pediococcus pentosaceus* (strain PPE1.0) DNA; Insertion sequence IS30 homolog (transposable element Insertion sequence IS30 homolog, kingdom Prokaryotae) DNA; *Pediococcus pentosaceus* (strain PPE1.0) DNA; Insertion sequence IS3 homolog (transposable element Insertion sequence IS3 homolog, kingdom Prokaryotae) DNA; and *Pediococcus pentosaceus* (strain PPE1.0) DNA.
ORGANISM *Pediococcus pentosaceus*
AUTHORS Leenhouts, K.J., Bolhuis, A.A., Kok, J.J. and Venema, G.G.
TITLE The sucrose and raffinose operons of *Pediococcus pentosaceus* PPE1.0
JOURNAL Unpublished (1994)
COMMENT On May 25, 1994 this sequence version replaced gi:475106.
FEATURES Location/Qualifiers
source 1..21838
/organism="Pediococcus pentosaceus"
/strain="PPE1.0"
/db_xref="taxon:1255"
repeat_unit 487..510
/note="DR2; putative"
/rpt_type=direct
repeat_unit 488..561
/note="IR1; putative"
/rpt_type=inverted
repeat_unit 513..561
/note="DR1; putative"
/rpt_type=direct
terminator 2724..2760
/standard_name="terminator 4"
/note="putative"
gene complement(2929..3864)
/gene="rafR"
CDS complement(2929..3762)
/gene="rafR"
/note="putative"
/codon_start=1
/transl_table=11
/function="regulation raffinose-operon"
/product="regulatory protein"
/db_xref="PID:g475107"
/translation="MNGEYKTLANKSFESNVLFQEA CLP NYTYKGNNVRDSYVIHY
IQEGKGTFAAANHPATVLKAGDIFILPKGTPCFYQADNDQPKWYFWIGFSAGIRIEAM
LSGSLLAQKCYLRQVQNGHIYADLSELYKVLHIPNSLINDVLLGSLIYRLFYDLLRWY
PADATNIKVKSTEQFNLAVSYLQENYSTGCTIMDLCHYLNLSRSYLYTLFKTHANTSP
QKLLTKRLEDAQRLLSTSNNSVQSIANMVGYKDSFTFSKAFKRYSGASPSYYRKSIG
I"
-10_signal complement(3837..3842)
/gene="rafR"

```

-35_signal      complement (3859..3864)
/gene="rafR"
gene            3909..5918
/gene="rafP"
-35_signal      3909..3914
/gene="rafP"
/note="putative"
-10_signal      3932..3937
/gene="rafP"
/note="putative"
CDS              3993..5918
/gene="rafP"
/note="putative"
/codon_start=1
/transl_table=11
/function="raffinose transport protein"
/product="permease"
/db_xref="PID:g475108"
/translation="MQEEHNYKWVGGRILIYGFAGKGNDAFYSILSGYLIIFITSHLFDTGNKALDNRMVSLVTLIIMVLRIVELFIDPFIGNAIDRTKNSPGHFRPWVVGGTVSSIIILLLFTNLGGLYAKNAMIYLVVFAILYITMDIFYSFKDVGFWMSMLPSLTDSREREKTATFARLGSTIGGGLVGVLVMPAVIFFSAKATSTGDNRGWFIFALIICLIALISAWGVGLGTREVDSDIRKKNQDTVGVMEIFKALAKNDQLLWAALAYLFYGVGINILGSLEVYFTYIMGKPKSFSILSIIINIFLGLIATSLFPVLSKKFSRKGVFAGCLVFMIGGIAIFTIAGSNLWLVLAAATMFGFPQQMVFLVVLMVITDSVEYGQLKLGHRDESLALSVRPLIDKFGGAISNGVVGQIAIISGMTTGATASSITAAGQQLHFKLTMFAPALMLLIAIGIFSKQIFLTEEKHAEIVAELETRWRTKFDNTTDQVAEKVVTSDLATPIAGQVIPLAQVNPDTFAAAGTLGDGFAIKPSDGRI LAPFDATVRQVFTTRHAVGLVGDNGIVLLIHIIGLGTVKLRGTGFISYVEEGQHQVQQGDELLEFWDPTIKQAGLDDTVIMTVNSTEFTMMMDWLVKPGQAVKATDNILQLHTKA"
gene            5976..8177
/gene="agaR"
CDS              5976..8177
/gene="agaR"
/codon_start=1
/transl_table=11
/evidence=experimental
/product="alpha-galactosidase"
/db_xref="PID:g475109"
/translation="MSLITVDQANRVFHLLNQTLSYIFAVEQGGTLSHLYFGGHVDHYHGELRYPRVDRGFSGNLPGSTDRTFSRDTLPKEYSTAGEMDYHLPAAIVRHTDGANALYLVYQGYRIEAGKPKLSQLPAAFVEDETEAETLTIVLVDQVSQVEFDLQYTIYRDRPVVTRSVQVCNQGDHVNLEKVASMQIDFTDQFETITLPGAHANERHPERGSINYGIQTFGSLRGTSQHMQNPFLALVDHTTTEFSGDAYGFNLVYSGNHALEKDQLDQLHLMVGINSYFNWQWKAGATFQTPEVLMVYTNKGLNAMSQAYHHLIREVRVVRSEFKNQERPIVVNNWEATFFDFNEAKLKPIVDEAKQLGIEMFVLLDGWFGHRDDDNSSLGDWQVDHDKFPOGLNHFVKVYHEQGLKFGIWLEPEMISYDSKLYQQHPDYLMQVPGRSPSPSRNQYILDLGROQAVRNNIFDQLDQLLKSQIDYIKWDMNRHLSDIYSVALPPERQGEVYHRYVLGLYELLERLTTAYPHILFEGCGGGGRFDAGMAYYMPQIWASDNTDAVARLTIQYGTSLAYPISLATAHSVSPNQQTGRETSMSTRSAVAASGVLGYEELDTQLSSADKQIVQKQVVOYKQIRPLIQFGEFYRLKSPITSNQAAWMFVSPQQDEAIVMVFNLTSYAQPSLTKTKLVGLNPKLNYQNIATKAIFGGDELMQLGFYDPVVYQDYTTKVYHFKAVTEN"
terminator      8265..8289
/standard_name="terminator 3"
/note="putative"
repeat_unit     9403..9426
/note="DR2; putative"
/rpt_type=direct
repeat_unit     10396..10442
/note="DR1; putative"
/rpt_type=direct
repeat_unit     10397..10443
/note="IR2; putative"
/rpt_type=inverted
terminator      10681..10716

```

```

/standard_name="terminator 2"
/note="putative"
gene complement(10727..11593)
/gene="scrK"
CDS complement(10727..11593)
/gene="scrK"
/note="putative"
/codon_start=1
/transl_table=11
/function="fructose 6-phosphotransferase"
/product="fructokinase"
/db_xref="PID:g475110"
/translation="MLLGAIIEAGGTFVCACTGAENGQVSDRISIPTTPVETMTAVDD
YFTTHPVDAIGIGSFGPIGVNPNDPKYGYITTPKPGWDFDFLGHLSQFNIPLYWT
TDVNEAAYGESMIGIAKDVPNSIYMTIGTGVGAGVISQNHIFNGRTHTELGHMRLNRL
PGDDFKSNCPCYHDICLEGIAAGPAVGKRTGKAGKDI PVDDPVWPIITDYIAQACVNLT
VAFAPDKIILNGGMNQRQLFPMIREKFAAYLNGYEEVPPLDDYIVPAGLGNNSGIAG
GLLLAQAALKNA"
gene complement(11673..13832)
/gene="agaS"
CDS complement(11673..13832)
/gene="agaS"
/note="putative"
/codon_start=1
/transl_table=11
/product="alpha-galactosidase"
/db_xref="PID:g475111"
/translation="MPVEYDPKTGLINLNHNDQISYVIQILAHRYPVHRYFGRYFSKQP
YFEPMPSGSHAFANDPTERFPYSVTSPLYEYSTIGSGDYRQPAYVIKDANNQLLPILE
YTGFSVNDQPINRQLPPTVSKHTPVTTLVIHLTDAVTKLQMDLNYTIFENQPLILRS
TTLRHGTTNLQVTALSSAQQLDPLTDQYTALTLSGTHAHEANPSFNRLHPGLQTVRSL
RGTSGPQHQPFMQLAE PNTTELAGTVIGCALAWSGNFDSTVEVDQYQHSRLTIGLEPD
TFEWQLKPNSSFQTPEAVLTWTNTGFNGMSQVFHDFSYQLMPSQTNIPSVLNTWETLT
FAVSESKVQHLLIEHAHQLGLQMLVLDDGWFVNNGENGQLGDWFVDPKFPNGLNPLA
QOAHHHRMKFGFWVEPEMITNSQLYQOHPDWLQYVDRTPITARHQVLQLDLSQAAVR
DHLITLTNLVQNNQOLDYLKWDMMRHLTQVGSTHLPAQQQGELYHRYVCGLYDILTRL
KRACPKLIIECSAGGGRFDFGMLPYTNQTWISDLTDPVDRATIENGFSYLFPPRIFS
NHITASPNAQNGRITPFETRLQLACIGQLGLELNPQQLAPSEQQLLRGALIKYQQLKS
TFIKAHFYRLPTTRHVVAVLIVTADKKQAIICCYLNGLNSRVKTQHPLPLHYLDALAY
SDSSGNRTGHQLNTMGIPLKPTNADFTSQLIYLCQN"
gene complement(13852..15989)
/gene="scrA"
CDS complement(13852..15807)
/gene="scrA"
/note="enzyme IIabc"
/codon_start=1
/transl_table=11
/function="enzyme II of the PTS system, sucrose specific"
/evidence=experimental
/db_xref="PID:g475112"
/translation="MNHQEVARVLNAIGKNNIQAAAHCATRLRLVIKDESKIDQQAL
DDDADVKGTFETNGQYQIIIGPGDVKYDALIVKTGLKEVTPDDIKAVAAAGQNKNP
LMDFLKVLSDIFIFIPIVPAVAGGLLMLNNVLTAEHLFMAKSVVEVYPGLKGIAEMIN
AMASAPFTFLPILLGFSATKRGFGNPyLGATGMGMIVLPSLVNGYSVATTMAAGKMVY
WNVFGHLVAQAGYQGQVLPVLGVAFILATLEKFFHKHIGKAFDFTFTPMAIVITGFL
TFTIVGPVLRTVSDALTNGLVLGYNSTGWIGMGI FGLLSSAIVITGLHQTFFPAIETQL
LANVAKTGGSFIFPVASMANIGQGAATLAIFFATKSQKQALTSSAGVSALLGITEPA
IFGVNLKMKFPFVFAAIASGIAASFGLFLHVLSVAMGPASVIGFISIASKSIPAFMLS
AVISFVVAIFIPTFIYAKRTLGDDRDQVSPAPSTVINVNDEIIISAPVTGASESLKQV
NDQVFSAEIMKGAAIVPSSDQVVA PADGVITVTDHHAYGIKTTAGAEILIHGLD
TVNLNGEHTTNVQKGDTVHQGDLLGTFDIAALKAAANYDPTVMLIVTNTANYANVERL
KVTNVQAGEQLVALTAPAAASSVAATT"
-10_signal complement(15872..15877)
/gene="scrA"
-35_signal complement(15894..15989)

```

```

/gene="scrA"
-35_signal 15993..15998
/gene="scrB"
gene 15993..17568
/gene="scrB"
-10_signal 16017..16022
/gene="scrB"
CDS 16063..17568
/gene="scrB"
/standard_name="sucrose-6-phosphate hydrolase"
/codon_start=1
/transl_table=11
/evidence=experimental
/product="sucrase"
/db_xref="PID:g475113"
/translation="MIWNRKTRYTPYEQWPATKLPQLVAQAROSKWRMQHHIQPTSGL
LNDPNGFSYFDGQWHLFYQVFPGPVHGLKSWQHVTSKNLVDWHDEGLAIRPDTPYDS
HGAYTGTALPIDDQLFIMYTGNVRTADWQRESYQLGAWMDTDNHIKKLSRPLIAHAPA
GYTSSFRDPDLIRNDHGYYALIGAQTTEIGAILVYFSKDLTTWTCQGELENVPANARG
YMIECPKSGLDRSTARLIVLSQGLSQATIPYQNIYPNMYLADQLNLAQAQFTEPHAL
TQLDDGFDVYATQAINAPDGRALAVSWIGLPEISYPTDRENWAHCLS LVKELTLDGH
LYQNPVAAVDDLRTTAHDLVFEQQRATVAALNGSFELLTVPADKTVTVNIADQQESG
QLQVTVDANHGQVMIDRRHTGNSFAEDYQGTRQVELTAHKTIKIRLIIDVSVFECYID
NGYSVMTGRFFLNATPSRLNVQGDITAVTGVVWEWRQSEHTGVDNNETKIK"
gene 17549..18529
/gene="scrR"
CDS 17549..18529
/gene="scrR"
/note="putative"
/codon_start=1
/transl_table=11
/function="regulator of the sucrose operon"
/product="regulatory protein"
/db_xref="PID:g475114"
/translation="MKPKLNDVAKLAGVSATTSRVINVNHGYLSSQTKEVFAAMREL
HYQPNNMARS1QGKNTRLIGVIFSDISNPFFGELVSRIEKILFAKNYKVILCNSADDP
QKERDYLQMLMANQVDGI1AGAHNLGIEEYQQYGLPIISFDRYLSDNIPIVSSDNYQG
GWLATQTLHQAGATNVAIFTGKSHAGSPTNGRREGYEAYLTAQQLTPHHELPFELTP
ALKMMEIKTIMTQHQYDGIFCSDDLALLVLNVAAQQLSLTVPEQLRVVGYDGTALIRD
YHSELTTEQPLADISTLLVSLLLQRIEDANCTLESKYTLPVKLIKGFTA"
gene 18550..20223
/gene="agl"
CDS 18550..20223
/gene="agl"
/codon_start=1
/transl_table=11
/evidence=experimental
/product="alpha-glucosidase"
/db_xref="PID:g493182"
/translation="MAATIKWWQQAVVYQVYPRSFQDTNHDGIGDLKGITAHLDDYLKQ
LGIDVIWLNPYIYRSPNDDNGYDISDYQQIAADFGTMADFDLLOAAHTRGLKIIIMDLV
VNHTSDEHPRFKRSRQDRTNQYRDFYFWRSGNGKKAPNNWEAAFGGSAWQYDEQTOQY
YLHTFSTKQPDLNWENPTLRESVTMMTWWLNKGVDGFRMDVINQISKLPGLPDGPLK
PHSQFGDARVTNGPRVHEFLQEMNQEVLSQFDIMTVGETHGVTPADALKYAGADQHEL
DMVFEFOHLLRLDNSQHGLGWSTRKTPVALKKVISDWQVGLEGRAWNSLFWNNHDTP
RAVSRFGDDRPAYRVSAKMLATCLHLLQGTPYIYQGEELGMTDAHFTELASYRDIES
LSAYRDLVTERQLLSPADMMARLAAASRDNSRTPMQWDTEVNAGFSDAAPWLTVPNPY
RQINAAAALADPDSVWYYQHLIQLRHQYPSVTLGSFELLWADDPQYSYMHNGKADL
ASLLQFHSDRTVPTTGSISDPTAKCLISNYGEQQPNKLRPYEAWVYQLA"
terminator 20341..20389
/standard_name="terminator 1"
/note="putative"
BASE COUNT 6181 a 4373 c 5015 g 6269 t
ORIGIN
1 gcatgcttga ctgggtgctt agccgcaa at gatcaggaga gaagcatcgc gactaggaaa

```

61 ggagggcaac acgggttgctt taatcaacaa agaccagttc gatagttca tcttcgtt
121 ttagtggac agataaatga cgattaaggt cgccaccag ttgcgtgcg aaggccatct
181 ttcgtcgaa ttccgtcatg tgcgaagtaa acggacgct ttgtaatta aagtttcgc
241 cgatatacgcc cgatagtc acgtgcga agtttttc gtcgtggcgaa ctggccgggtt
301 tcaggacggt atccaaatac tgatagagaa tcgtggatc gtagttcagc ggcttatcaa
361 gggtcggtt aagctgttag ccgggtcgaa gaccagcgaa ttgttcgtt atttagactt
421 ggtaaggct ttgatacaat ttcaaaacga tcactcctct tcttttcaact tttagatag
481 cagttatggt agattgtaaa attaataccga acgtgtatc ttttgcgg aacagcggtt
541 ggattaattt tacaatctac cttttacca gcaattgtatc aaatgctagc taacgctagt
601 cctagtata aattgtctaa acgcttgcgaa gcaatcttgc cccaaagcatc tatttctcg
661 cgtatgtt tgggtcaat gattggcggt gtatgaatatttgcgtacact aaaaatgtat
721 catttgcagt taatgaattt aacacagcaat atatcgatgg ctataataat ttaagcccta
781 ataaaactatc aacagttcca gataaagtag ccaaccaagc tgtagtatttca
841 ggggtcgatgg cgacgtcatg acgacacgaa taaaagaaac cctgaacaag ggtttgagtc
901 gcccgtgaa aaaatcagac atcaaaaaa tgaccgcgat tattcctcaa agcgggtatc
961 gaattgtatcaatttgcgaa acgcctatga accaaatgtatc atccagaatt catacgtaa
1021 caactaatga agctatacgc aatagcaacg gtggtaagaa tagaactatc
1081 atgttcgtatgatgtgg ttaaccggg aagattgtatc ttttttttttgcgtatc
1141 ccaacttgata gacaaacatttgcgtatgatc
1201 ccaagggtgtc gttccagct ggtagccgtt gatgaatgtatc
1261 cttgtataggg tatttaagga aaaaaaaaaa cccacttagt
1321 aatattatta attggcttcg ctcacaacaa agtcgagggg
1381 ataatttagt ataagaacct taaatgaagc taaaactatc
1441 agaaagaaac tacagcaaca acaattactg ctccaaataat
1501 cggctaattt agggcccaa gatccgagta tagttctcc
1561 cggcaacaat attagcaatc cacccttgcgtatc
1621 tagcaccaat gatggcacca ataattaaaa cccataacca
1681 tcttattttt cgagggatgt ttttttagatt tgttatagtt
1741 ctaataaaag ttgagatata gaaatgataa ttgataattt
1801 tcatatgaaa tcctccattt tctaaatgcgtatc
1861 atatttgcat acaagttatttgcgtatc
1921 ctttttctc aattacagac gttaaagaat aattgattt
1981 ttagtctac ggacttaaaa aggagtttgcgtatc
2041 gatcctaacc cagaacctaa gccaaacatc
2101 gcgtgggtgt atagacggtt
2161 gaatttcaat ggtacctgaa tttttcgatc
2221 tggcttaactt aattataaaa cgcgcgagaa taaatttttgcgtatc
2281 ccataataaa ttatttctat tgaatttttgcgtatc
2341 agtaactcaa gctagtttgcgtatc
2401 tgcgcctaaa gagatgcgtt
2461 tggaaaata cagccggat
2521 gaagagttca tgcagactt
2581 aagtgggtat gattgactaa
2641 taattaaata aacgcgatttgcgtatc
2701 ataaacaaa taatttttgcgtatc
2761 ttgtttaaa aaaggctgtatc
2821 gcgttattac ggtgttacca
2881 cattagcaga aagacaaatc
2941 gcttttacga taatagctgg
3001 gaagctatcc ttgttagccta
3061 tagacgctgt tttagcatctt
3121 agcgtgcgtt taaaacaagg
3181 gtccataatg gtacaaccatc
3241 ttgttcgtt gactttactt
3301 atagaacaat cgatagatca
3361 gtgtaaaact ttataaagtt
3421 taagttagcat ttcttagtgcgtatc
3481 gggaaaacccg atccaaaatc
3541 tggccctta ggtgtatc
3601 tgcagcaaaatg gtacccatttgcgtatc
3661 attaccttgcgtatc
3721 ttgcgtatc ttattcgtatc
3781 taaaatataat tgctaaatat
3841 tactgttatt taaaacatgcgtatc
3901 qattactatg ggcaacccaa

3961 acaaaaatga tttgagggca ggtaattcat ttatgcagga agaacataat tataaatggg
4021 taggaggtcg cctaatttat gtttttgggt caaaaggtaa tgatgcattc tatagtatcc
4081 tctccggata ttttaattatc ttcattactt cacaccttt tgataccggc aacaaggcac
4141 tagataatcg aatggtcagt ttggtagct tgattattat ggtgttacga atcgttgac
4201 tattcattga tccctttatc ggtaacgcaa ttgaccgaac taaaaactcg ccgggtcatt
4261 tccggccatg ggtttagtc ggtgggaccg tctcttcaat tattctctt ctattattta
4321 ctaatttagg cggccttat gctaaaacg caatgattt tttggtcgtt tttgcaattt
4381 tgtatattac gatggatatt ttctattcg taaaagacgt tgggttttgg tcaatgttac
4441 ctcccttgac cactgattca cgagaacgcg aaaaaactgc tactttgcc cgcttaggg
4501 caactattgg tggtgctt gttgtgtgt tggtcatgcc ggcagttatc ttctttctg
4561 ccaaggcgac tagtactggc gataatcggt gttggttcat tttgcctt atcatctgct
4621 tgatagctt aatttcagcc tgggggtgtt gcttaggtac acgtgaagtt gatagtgata
4681 ttctgtaaaaaa taaacaagat acagttggg tgatggagat ttttaaggca ctggctaaaa
4741 atgatcagtt gctatggca gcttagctt atttattcta tgggtgtggc attaataattt
4801 taggttcaact ggaagtctat tatttcacgt atattatggg taagccaaaa tcattttcta
4861 ttctatcgat cattaatatt ttcttaggtt tgattgtac atcgctattt ccagtattat
4921 cgaaaaaattt cagtcgtaaa ggtgtttttt ctggctgtt agtggtcatg ctaggtggta
4981 ttgcgattttt taccattgtc ggcagtaatt tatggctgtt tctattagcg gcaactatgt
5041 ttgggttccc gcaacagatg gtcttttag tagtttgat ggtatcacg gattctgttg
5101 aatacggtca attgaagctg gggcatcgat atgagtcgtt ggctttatca gtgcggccgt
5161 tgatcgataa atttggtggc gctatctca acgggggtgt tggccaaattt gccattattt
5221 ctggatgac gactggtgcg actgttttctt ctatcactgc tgcaggacaa ttacatttt
5281 aattaacgt gtttgcctt cccgcgttaa tgctacttat cgctatttgc atttttcaa
5341 agcaaataattt cttaccgaa gaaaaacacg ccgaaattgt ggctgaactt gaaagaacct
5401 ggagaacaaa gttcgataat accactgatc aggttagcaga aaaagtagtt acatcaactt
5461 atttagctac accaatcgct gggcaagtga ttccacttgc ccaagtcaat gatccaactt
5521 ttgcggctgg aacgttaggt gacggatttg ctattaaacc tagtgatggc cgaatatttag
5581 ctccatttga tgcaacggta cgtcaagtat ttaccacacg acatgcagtt ggcttagtcg
5641 gtgataatgg gatcgcttta ttgatccata ttgggtttggg aactgttaaa cttagaggaa
5701 cgggattttt ttcttatgtt gaggaggggc agcatgtaca acaaggggat gaattacttgc
5761 agtttggga tccaaacgatc aaacaagctg gtttagatga tacggttattt atgacagtga
5821 ctaattcaac cgaatttact atgatggatt gtttagtcaa gccaggtcaa gccgttaaag
5881 caactgataa tattttacag ttgcataacta aagcataatg tgccactggt tatttttcaa
5941 tttaaaattt agtaattccc cgaggaggaa ataaaatgtc attaattacg gttgatcaag
6001 cgaatcggtt tttcatttgc cataatcaaa ctctttcttgc tatctttgc gttgagcagg
6061 gcggtacttt aagtcatttgc tattttggg ggcattgtggc ccattaccac ggtgaatttgc
6121 gctatccacg agtcgatcgt gtttttccg gtaatttacc gggatcgact gatcgcaactt
6181 tttcacgtga tactttgcg aaagaataaca gtacggctgg taaaaatggat tatttttgc
6241 cagcagcaat cgttcgatc actgtatggg ctaatgcattt atacttagtt tatttttgc
6301 atcggtatcga agcgggtaaa ccaaaggtaa gcggttacc ggcagctttt gtggaggatg
6361 aaacagaggc agaaactttg actatcgat tagttgatca agtaaggccaa gtcgagtttgc
6421 atctacaata tacgatctac cgtgatcgac cagttgtcac ggcgttgcg caagtctgt
6481 atcaagggtga tcatgctgtt aatttagaaa aagttgttcc aatgcataatc gatttttgc
6541 atcggtatcga cggaaacattt acattaccgt gaggcgatgc caatgaacgc catccagagc
6601 ggggttcgat caattatggc atacagactt ttggtagttt gcccggactt tccagtcatt
6661 aatgaatcc atttttggcc ttgggtgtatc acactacgac agagtttagt ggtgatgcatt
6721 atgggtttaa cttggatcatt tctggcaatc acgcatttgc actggaaaag gatcaacttgc
6781 accaacttca tctaattggc ggttataata gttataattt caattggcaatc cttaagctg
6841 ggcgtactt ccaaacacca gaagtataa ttggcttatac aaataaaagg ttgaacgcga
6901 tgagccaaatc ctaccaccat ttgatccgtt aacgagtagt acgttagttagt tttaaaatc
6961 aagagcgatcc gatttgcgtt aacaactggg aagcaacattt ctttgatattt aacgaagcaa
7021 aatttgcgtt gatcgatggat gaaaggccaaatc agttaggcatt taaaatgtttt gtttttagat
7081 atgggtttttt tggacatcgt gatgtatcattt attcttcattt aggtgacttgc caagttgtatc
7141 atagaaaaattt tcccgaggcc ttaaatcattt ttgtgaaata tggatgcataa caaggccatt
7201 aatttgcgtt ttgggttagaa ccagaaatgc tttcatatgc ttcaaaatgc tatcagcaac
7261 atccagatca ttaatgcattt gttccctggcc gatcaccatcg tccatcactt aatcgttata
7321 ttttagatctt tggatcgtt gatgtatcattt tgaccatcg tggatgcattt gatcaacttgc
7381 taaaatcgaa gcaatttgcgtt tatattaaat gggatgcataa tggatgcattt tcaatcgatatt
7441 attccgttgc ttaccacca gaaacggccagg gtggatgttcaatcgatgtt gtttttaggt
7501 tgtatgttattt atttgcgtt gatcgatggc cttatccgcgca tatctttttt gaaagggttgc
7561 caggtgggtgg cggatcgatcgtt gatgtatcgtt gttggatgcataa tggatgcattt gatcaacttgc
7621 gtgataacac tggatgcgtt gatgtatcgtt gatgtatcgtt tggatgcattt tggatgcattt
7681 caatttcgtt agtaccggcc catgtctcgat tgagccctaa tggatgcattt gatcaacttgc
7741 cgtcaatgtc gactagaatg gcaatgcgtt gttggatgcataa tggatgcattt gatcaacttgc
7801 taacacaactt cagttcgtt gataaacaatc tagtccaaatc gcaatgttgc tggatgcattt

7861 agattagacc actgattcaa tttggtaat tttatcgctt aaaatcacca attactagca
7921 atcaaggcgc atggatgtt gttcaccc aacaagatga agcgattgtc atggatattt
7981 atctgacatc ctatgccaa cccagttga caaaaaccaa attagtcgg ttaaatccca
8041 agctgaatta cccaaatatt gctactaagg cgatattgg tggatgtgaa ttgatgcaac
8101 ttggcttcta tgatccagta gtctaccaag attacacaac aaaggtctat cattttaaag
8161 cagtcactga gaactaactt ttacgtaaat tgcaatagta agttaaccag tgccggtaaa
8221 gtacacagac gaaaggcgc atgggagtc cagggtgatc aggccgcgc gccttaccc
8281 gcacgcgcta attccttcg aaatacttca gctgggtac agaacctaag taaacggcgc
8341 gggagatggt tgagccgaga ttcaagttga cgaaccagac tgggtgtac tagatcgagt
8401 gattaccct tagaaagta acgtctgagt gattgtggac ctcattagtg gcgcttcgg
8461 ccgagctata agggtggta taataagttc cgtaataacc ctcagcgca gccgttagcg
8521 tggtaactc tggtccgtt tccggctaa ctgtccgcatt aacgttaccg aattcagccg
8581 tagtattttg catgcataag ccacagatg cgcgtttac gttcggtaaa cgtaatagg
8641 acactctcg taccaggcagc ttggccatg acagatatcg tctcaaaatg accaaactcg
8701 tgacgcattt ctactcgctt ttggccgttca tcaataactac gaccgaatag ggcgtatgt
8761 tggtgcgagt ggtgtgggt catccgacgt tttgtttt ccagcagatc aaggtaacgg
8821 atctccaaca gttgcgcattc gatatagttac tacaatgttca tcgtacagac cattttggcc
8881 ggcttaataa accgggtcgtt acgggcataa ccaacggcag cgtccggca ccagccctca
8941 tggcagaact tatcatcaaa gttaggctaaag aaggccgtca cttggctaaa tttaatggg
9001 cgtccgcagt ttggccgtt tatttcatacg cgattctgag cggcttcagg actgttaatt
9061 tcaaagtatg gtttctgatt attaacgcgt ttaacttgcg tgactcgacc acgcttgac
9121 tcattattga cggtctgggt gcaagacgcatt aatcaatcg ctatttgcg tactgagcat
9181 ctttcaatgt gccaagcggc gatttaaccc cggtcgattt ccgataaaatg gtgacccccc
9241 aggcgtgggt tgatagactg ttcttgcattc aagacgaaaa cttcttccat tgggtttgt
9301 aggaacttca atgatacctg aattttctgt cttgggtgtct ttttttacca ttttggctgc
9361 gtggctaact taattataaa acgcgcggat aattttgata attggtagat tgcaatattt
9421 atccgaattt ttggacaaat ttgtcagcat aacctgatac ggtgtttgtc agtcgagat
9481 ttaagcggt cgctggtaa ttggagtaa cgtcgctgtt aatcttgcg tactaatgtg
9541 ctcaaaacga gtcctttag gataaaaaca acgttaattt cgttgcgtt gttcattact
9601 accacgtca gtcggatata agcatggcag taataggctc taataccata ttgtgattca
9661 agtgcatact gaccgctaaa ctcagtgcca cggccaccgc tgaagctgtg caccggacca
9721 ttaaaaagtgg ttaggaactt agttagggct tcattaaacag tcgcgtcgccgatcttt
9781 aaccggtagg cccaaaggaa ccgtgatttgcgatcgattt aagttaataa aactgcctta
9841 ctatgccac gaggaccaac gactgtatct agttcaaaat cggatggatcgttgcgatt
9901 ttaaacatca tgggacgctg ttcaattgtat cggcccaaaatg attgattata tttggatcg
9961 tggtaacgt tacggcgtt gcgtacgcca tggtaggttgc gatcattcaaa gggaaacca
10021 attctccctt gatttagcca attataaataa gatttagtag ttagtttttttgcgatt
10081 atcattcctg gtgaccagct tagacgtaaa tggtaggaa ttttttgcgatttgcgatt
10141 ctcagcttag tttccgacc acattgtgtat cggatgttgcgatttgcgatt
10201 tcagcctgat aagggtgaca tcgagataat tcataagaaa ttgttgcgg tgatcggtt
10261 agccgaacgc ccatttggat attggacacgc cctagttcac aaaaggttgcgatt
10321 ctttcggaaat aggttataact agacaaaaga tcagtccttca aaagatgggt ttgtggtaaa
10381 caccatttttta aggaagctga tcttttgcgat cggatgttgcgatttgcgatt
10441 ccatgtacgt gtcgtatatt ccgcaaatca tttcaattt ctcaggtgac ccagtatcg
10501 cgctacaacc actcggttgcgatcgttgcgatttgcgatttgcgatt
10561 agacttataa ggattggccc gtttattttt ctttcggatcggatgttgcgatt
10621 tcactgtttt aaccgtatatt atgcattttt cggatgttgcgatttgcgatt
10681 ccgctaatac gtttaaacac gtatcgccgg actttttgc aagggtttaa ggttcttta
10741 aagcgcctg agctaaacat aggccaccgg caatgcgatcgttgcgatt
10801 gaacgatata gtcatccagt ggtggactt ctttcggatcggatgttgcgatt
10861 ttccacgaaat catggaaag agtgcgttgcgatttgcgatt
10921 ttcacggtgc gaaggcgcg gttgggttgc ctttcggatcggatgttgcgatt
10981 ttggccagac cggatcgatcgttgcgatttgcgatt
11041 ctgctggccc agcagctaaat ctttcggatcggatgttgcgatt
11101 aatcgcccc aggttaaccgg ttggatcgatcgttgcgatt
11161 taaaaatgtt gttctgacta atgacgcgcg cggccaaatcc ggttccatcgttgcgatt
11221 tgctgttagg gacgtccttgcgatcgttgcgatttgcgatt
11281 catcgtgtt ccagtagat gggatgttgc ctttcggatcggatgttgcgatt
11341 agtcacccca gccgggttttgcgatcgttgcgatttgcgatt
11401 taacgcccgtt gggccaaat gaaccgatcgttgcgatttgcgatt
11461 agtcgtctac cgcagtcgttgcgatcgttgcgatttgcgatt
11521 taacttgcctt gtttcggatcgttgcgatttgcgatt
11581 cacctaaaatgatcgttgcgatcgttgcgatttgcgatt
11641 attcttaattt cggatccatcgttgcgatttgcgatt
11701 ctatgttgcgatcgttgcgatttgcgatttgcgatt

11761 gtatagcgat tacccgatga atcgctatac gctaattcag catcgagata gtgtaatgg
11821 aatgggtgct gagtttgac acgagaatta agcccattta agtaacagca aattgcctgc
11881 ttttatctg ccgttactat caaccacgc acaacgtgc ttgttggtagacgataa
11941 aagtgggcct tgataaaaagt agatttcagt tgctgatatt taattaaggc tcctcgaaga
12001 agttgctgtt cactaggagc aagttgctt ggattaagtt ccaatcctag ttgcccgtat
12061 caggctaatt gtaatcgat ttcaaatggt gttattcggc cattttgcgc attgggtgat
12121 gcggttataat ggttgcgtt gtaatcgat ttcaaatggt gggaaatagat aactaaatcc gttctcgat
12181 gttgctcggt cgactggatc agtcaaatacg cttatccacg tctgattgggt gtatggtagc
12241 atacccaaaat caaaccggcc accaccagct gaacaattt cgataatcag cttggggcag
12301 ggcgcgttca agcgtgttag gatatacataa agtccacaga catagcgatg atacagttcc
12361 ccctgttgcg cagccggtaa gtgtgtgcg ccaacctgtg taaggtggcg attcatatcc
12421 catttaagat aatcgatgtt gttattttga accaagttcg tcagagtagt gattaagtgg
12481 tcacggacag cggcctgcga aaggctcaag actagctgtt ggcgagctgt gatgggtgtc
12541 cgatcaacat attgtataac ccagtcaggta tggtagtttgggt agagctgact gtttagtgg
12601 atcatctctg gttccaccca gagaccgaat ttcattcgat gatgatggc ttgttggca
12661 aggggggttca gaccattagg aaacttaatg gggtaacaa accagtccacc taattgccc
12721 ttttcgcgtt tacgggtgac aaaccagccg tcgtctaaga caagcatgtg taagccta
12781 tggtaggggtt gttcgattaa gtgttgaact tttagactccg aaacggcaaa tgcgtgtt
12841 tcccaagtgt taagcacact tggatgtt gtcgtggagg gcatcagctg atagctaaag
12901 tcgtgaaata cttgcgacat gccgttaaaa ccagatattt tccaagtttag cacggcttca
12961 ggagtctgaa aactgtatt tgggttgagc tgccattcga atgtgtcggt ctcaagacc
13021 attgttaggc ggctatgtt atattggtca acttcgacag tactgtcgaa gttaccact
13081 catgctagtg cacaaccaat aacagtccca gctaattctg tcgtgttagg ttctgctaga
13141 gccatgaacg gttgggttt tggccacta gtgcgcgc gactgcac cggttgc
13201 cctgggtgtt aacggttaaa cgatggatc gttcatgtg catgggtacc gcttaatgtc
13261 agcgcgttat attggtcagt tggtagatct agttgagcgc tggataggc tggtagtt
13321 aaatttagtag taccgtgatc ggcagatgtt gtcgttcgcg agataagcgg ctgatttca
13381 aaaatagtgt agttaagatc catctgttac ttggtcactg catcggtcaa atgaatgact
13441 aggggtgtt caggcgtgtg tttgctaaacg gtgggtggca atggcgcga gttgattgg
13501 tgcatttca cagaaaaacc agtataattt aaaaattggca gtaattgatt gttagcatcc
13561 ttaataacgt aagctggctg gcgatagtca ccactgcgg tagtggataa ttccgtgg
13621 agtgaagtca cggaaatattgg aaagcgttct gttgggtcat tagcaaagc atggctacc
13681 gatggcatcg gctcaaagta cgggtttt gaaaagtaac gtccaaata tcgatgcact
13741 gggtaacgtt gagctaaaat ttgatataaca tagctaattt ggtcattatg aagattgatt
13801 aagccgggtt tgggtcata ttcaacaggc atgacgtcct cctaataaag ttttagacagt
13861 tggtagcgcgactgatgtt ctggccggc agtcaacgcg acgagctttt caccagctt
13921 gacattcgatc actttgatc gttcaacatt agcgttattt gccgttattt tgacgattag
13981 catgacgggtt ggatcatatg tagcggttt taaggcagcg atgtcaaaacg ttccgagtag
14041 atcaccttgc tgaacggat cgccttctg aacgttagt gtaagttt caccattaa
14101 attgaccgtt tctaattcata agtgaatcag aatttccgcg ccagcagg tttgtatg
14161 ataagcatga tggctatcat aagtgcacgt gatgaccccg tcagctggc cgacgac
14221 atcagagctt ggcacgattt cggccctt ccccatgatt tcggctgaga aaactgg
14281 gttacttgtt ttgagacttt cgttagcccc cgttaccggc gcaactgataa tctcatcatt
14341 aacattgtatc acggactctg ttgggtctgg tgattttact tgatcacat cgtcacc
14401 tggtagtttgc gcatagataa aagttggaaat aaatgcacg acgaacgaga tgacag
14461 gagcatgaat gctggatcg actttgaacg aatcgaaatg aatccgatga ctgatgcgg
14521 accccatcgcc acggacagaa cgtggatag tcctaagaaa gctgaagcaa ttccgtgg
14581 aatcgctgca aagacaaatg gaaatttcat ttggaggtt accccggaaaa tagcagg
14641 cgtatattccg agtaacgtt agacccggc tgaagaagtc agggctttt gctttgg
14701 ctttagtggca aagaaaatag ctaaagtggc ggcacccgtt ccaatgttgg ccatcgagg
14761 gactgggaag ataaaggaaac caccagttt ggcaacattt gccaacaact gggttcgat
14821 tgctggaaa gtttgcgtt gaccgttata aacaatttgcg gatataata aaccaaaag
14881 tcccatacca atccagccgg tgctgttata taaggctact aaaccgttag ttaatgc
14941 gctcaccgtt cgtatggatcg gaccaacatg tgtaaaatggt agggaaacc
15001 ggcaaaatc gggtaaaacg tgaagtcaaa tgcccttta atgtgtttt ggaagaattt
15061 ttcaagggtt gctagaatgcg aggcgcaccc taagactggg agcacttggc ttgtatag
15121 gcttgcgtt acgtgttacc caaagacgtt ccagtaaaacc atcttgcgg ctgcgtt
15181 ctttagtggca cttatggatcg taacttagtgc tgtaagacc atgatcatgc
15241 accccatcgcc acgggttgcac cgaacgcgtt cgttgcgtt gatcataa ggttgg
15301 gaaagtaaaac ggcgcactatg ccatcgccgtt aatcatttgcg gcgatcactt
15361 gtaaaacttca acaactgact tcgcccataaa aagatgcgt gcaatcgggg
15421 tgccattaaat agaccccccgg caacttagtgc agggacgatt ggaataaaaa
15481 gactttgggg aagttccattt atggattttt attttgcacc gcaacgcgtt
15541 gtcatcggtt gtcacttctt taagacccgtt tttaacgcatt aaggcgtcat
15601 gacatcgccca gggccataa tgatttggta ctggccgtt gtttgcagg tccccttaac

15661 gtccgcgtcg tcatctaagg cttgttgatc aatcttgat tcatccttga tgaccaaacg
15721 gaggcgtgtc gcacagtggg cggcggcttg aatgttggtt ttaccaattt cattaagtac
15781 gcggcggca acttcttgc gattcataat aattcactcc ctgagtctaa aattatttt
15841 aatttaataa tgcaagcgt ttacagaccc aataatagca attttggaaa atatgtcaag
15901 cgtttgatac aaaattggat tgcttgataa tctggttaaa aagccgaatt tgaaacgctg
15961 acatgaaaat ataataaaa aatgtcaaac gattgacata tcggtgtaa cggtttata
16021 atagtgacaa ttaagtgcga cgttaaaag gagctcatcg tcatgatatg gaatcgtaaa
16081 acccgttata ccccttatga acagtggcca gcaactaagc taccacagct cggtgcccag
16141 gcacgccagt ctaagtggcg gatgcaacat catattcagc ctacatcagg ttgtttaat
16201 gaccccaatg gctttcata ctttgatgtt cagtggcacc tggcttacca agtcttccg
16261 ttccgtccgg tacacggact gaagtcatgg caacacgtga cctctaagaa ctgggtggat
16321 tggcatgtat agggcttagc gattcgcacca gatacggcg acgactcaca cggcgcgtac
16381 acgggtacag cactaccaat tgatgatcag ttatttatta tgtatacggg taatgtccgt
16441 actgcccact ggcaacgtga atcgtatcaa ttagtgcct ggatggacac ggataatcac
16501 attaaaaaagg tggccggcc actaattgcg catgctccgg ctgggtacac gagttcggtt
16561 cgtgatccgg acttgcgttccg gaatgaccac ggctactacg cgctgatcg tgcgcagacg
16621 actactgaga tcgggtctat cttggtttat ttctcaaaagg atctgaccac ctggacgtgt
16681 caaggcgagt tgaacgttcc cgcaaatgcg cgggttata tgattgagtg tccaaaatct
16741 ggtctggatc gatcaacagc cggcttattt gttctgtccc aagggttatac gcaagcaacg
16801 atcccgatcc agaatattt tccgaacatg tacttagttt ccgatcaattt gaaacctagct
16861 caggcacaat tcactgaacc ccacgcctt acgcaattgg atgacggctt tgacgtgtat
16921 gccacgcagg caatcaatgc gccggatggc cgccgcgttag ccgtcagctg gattgggtt
16981 cctgaaattt cctatccaac tgatcgtgaa aattgggccc attgtttagt ctagttaag
17041 gaactaacgc taaaagatgg gcacccctat caaaatccag tggcagcggg ggatgattt
17101 cggacgacgg cgcacgtaccc aactttttagt caacaacggg cgacggcggc cgctttgaat
17161 ggcagcttt aactcttact gacgggttccc gctgataaaa cggtcacggg taacattgct
17221 gatcagcagg aaagtggcca gtcgcaggactg actgttgcgtt ccaaccatgg tcaggtttag
17281 attgatccgg ggcatacggg gaattcggtt gctgaagatt atggtcagac tcgtcaagtt
17341 gagttaacgg cccataagac gataaaaatc cggctaataca tagatgtgtc agtctttgaa
17401 tgctatattt ataacggcta ttccgtgtat accggccgtt tcttcttaaa cgctaccccg
17461 agtcgtctta acgttcaagg tgacactacg gcaggtaaccg gtaaagtctg ggaatggcgc
17521 caatcagacg atactggagt cgataacaat gaaacaaaaa taaaatgatg ttgcgaagct
17581 agcgggggtc tcagcgcacca ccgttctcg gtttattaaat aaccatggct acctgagtag
17641 tcagactaaa gaaaaagttt tcgcggcgat gcgagaactg cattatcagc cgataataat
17701 ggcacgctca ctgcaaggaa agaatacgcg cttatttgcgtt gtcattttt cagatattag
17761 taatccgttc tttggtaac tgggtcccg gattaaaaaa atcttattt ctaaaaattt
17821 taaggtgatt ttatgtataa gtgcggatga tccgcaaaaaa gaacgcgact attacaaaat
17881 gttgatggct aatcagggtcg acgggattat tgctgggtcc cacaatttag gtattgaaga
17941 atatcaacaa tatggtcttc cgattatttc atttgcacgg tacttattccg acaatattcc
18001 catcgtagt tcggataattt accagggggg ctggctagca acccagacat tgcacgcggc
18061 gggggcgact aacgtggcta tttttactgg taagtccac gctggttcgc caacgaatgg
18121 tcgcgtgaa ggatatgaag cctacttgc tggccaaacag ctaacacccc atgtccatga
18181 gttaccgtt gaattgaccc ctgcgtgaa gatgatggaa attaagacga ttatgacaca
18241 acatcagttt gatggtattt tttgttagtga tgatttggcc gctttacttag tgctcaacgt
18301 cgcgcacaa ctgtcattaa cggttctga gcaattacgc gttgttgggtt atgatggcac
18361 ggcgcgtata cgtgattatc attctgaatt aacaacagta gaacaaccgt tagcggatata
18421 tagcacgtt ctgggttctc tactatttgc gcggttggag gatgcgaattt gtacactgg
18481 atcaaaatat acgttaccgg ttaagctcat caagggtttt accgcgttaac taaggagtt
18541 atcgtatgg tggcgcacaa aatcaagtgg tggcaacaag ccgttgcata tcaggtctat
18601 ccacgggat ttcagatcatc caatcatgtat ggttattggc actttaaagg gatcagggcc
18661 cacctagact actttaaaggca attaggaatc gacgtcattt ggctcaatcc catttatcg
18721 tcacccaacg atgataatgg ttatgtatattt agtggatttcc aacaatttgc gggcacttt
18781 ggcactatgg ctgattttga tgaactattt caagccggcc atgaccgggg cttaaagatc
18841 atcatggact tgggtgtttaa ccataccctt gatgagcatc cttagttcaa acgcagtcgc
18901 caggaccgca ccaatcaata tcgtgttcc tactttggc gctcaggtaa cgccaaaaag
18961 ggcaccaata attggggaggc ggcttttgg ggttcagcat ggcaatatga tgagcagacc
19021 cagcgtattt atttacacac gtttcaaca aaacaacctg atttaaactg ggagaatccg
19081 accttacgag agtcaattt tacgtatgtat acctgggtggc tgaacaaggg tgggtacgg
19141 tttcaatgg acgtcattaa ccaatctt aagttaccgc gattaccaga tggccactg
19201 aaaccgcaca gtcagttgg cgatgctcg gtaacgaacg gaccgcgact tcatgaattc
19261 ttgcgtttaa tgaaccaaga agtattgtca cagtttgata tcatgacggt tgggtacgg
19321 cacgggggtga caccacgcga tgcgtgttcaag tatgcggccg ctgatcagca cgaatttagat
19381 atggtctttt aatttcaaca tctgcgtactt gataatagtc agcatggct tggtaagtgg
19441 agtacgcgtt agacaccgtt agtggcgtt aagaaagtca ttagtgcactg gcaagtcgg
19501 cttgaaggc gggcttggaa cagcctgttt tggacaatc atgatacgcc acgagcggc

19561 tcacgtttg gtgatgaccg accagcatat cgggtacgtt cggctaaaat gctcgcaacg
19621 tggttacacc tcttacaggg aaccccatat atttaccaag gtgaggaact ggggatgact
19681 gacgctcatt ttactgaact agccagttac cgagatattg aatcgctcag tgcctatcgt
19741 gatttagtga ctgagcgcca gctgctatca ccggcagata ttagtggcccg attagcggcg
19801 gcttctcggg acaactcgcg aacgcccgtg caatggaca cggaggttaa cgcgggttt
19861 agtgacgcag cgccatggct cacggtaat cccaaactatc gtc当地atcaa tgctgcccga
19921 gcttggctg atccggattc agtgtggta tattatcgc acttgattca attacgacat
19981 caatacccct cagtgacgtt aggttcgtt gagctctgt gggccgtatc tcccgagtat
20041 tcttatatgc acggcaatgg gaaggcggac ctggctagtt tggtcaatt tcacagccga
20101 gacactgtcc cgaccactgg atcaatatct gaccccacgg ccaaactgtt gatttagtaat
20161 tatggcgacg agcagccaaa taaattacgg ctttatgagg cttgggtcta ccaactcgca
20221 taggtattca ggacttggct aattgaggtt gtcttggaa tatacccaat cgttacgatt
20281 agtgtgaatt gggttgttca aaattctgcg atggcttaa taagcgttca cctgactaaa
20341 aaaactgttt tcaaaaaggat aacttttggaa agcagttttt ttagccggg cgaccacttg
20401 gcccacgtga ggctcatcta gtcgttggg accttgaatg aagcccgatc ggttaaacgg
20461 tagttgttca tcgatcgatc tgaccaattt accagttatct agcaactgtt atagtaattt
20521 taaggcggct tgatcagaaa tcaactcgat cgggtggatt gcttggaaac gaattgattt
20581 actgttggct ggcaagtgttca gggcgaatgtt aagcgttggc gcatcaatt tggccattgc
20641 taagtcatct ttgtccaaac gccatcgatc gtcgttggg accttgaatg aagcccgatc
20701 gggcttggat gtaagctgtt ccctgagaca actttttaaga gagggtataa tgaataatc
20761 gctctcaaa agaaaggaat ttttacctgc caactcgat cgggtggat gcatcaatt tggccattgc
20821 acattatcaa cctatataag caaggcgtt cagctggccca actggccaga gaatatggca
20881 ttggcttattt aacagttcat aagtggatcc agggccaggc caaaactcaa tccggtaat
20941 cgccagacga aatcaaagcg atggaaaagc gacgggcttc gctgtctgat gagaacgaaa
21001 tcctaaaaaaa gcccgggtt tccttgcgcgaa gaaatcaaccat atatcttgc ttacgttcc
21061 caagaagccca tcaccaccat gtaaccaaga tggccgtt cctcggtt tccagagctc
21121 agtattatcg ttatcgatcc cccaaacccat caaaacggcc gggcgaatgtt gcccggat
21181 aacaacggat tctcggttcc ttggggat ttaagcgttca atacgggtt atgaagatcc
21241 accatgaattt gaaatcgatcc cttcaaccatc tgcgttcc gtcgttcc caataagtgg aaagccggctt
21301 cccggctcat gaaaggactt gatatccact ccgttaccgtt taagcgttcc ttctcgacca
21361 cggcttccaa aaccaaggat gaaacgcgtt ccaacttgc tcaaaacggat gcccggat
21421 ctggtttaaa tcaaaaatgg accgctgatc tgacccat tcaaaacggat cgtatggct
21481 ggtgttactt atcaaccatc atggacctgc actcacgtt gattatccg tattcggtt
21541 caaaaaagat ggctactgtt ttagtctttaa agaccctggaa aagcgttcc aaaaatcgaa
21601 ccattactgg tggacctgtt tatccatatacg gatttaggtt cacaggatac cagcgattgt
21661 ttacaatcaa cgtttaactt agctacatcc cccgactca tacagccgtt aggttatcc
21721 tatgataatg cgccaaataga atcccttcac gcttccctca aaaaggatg tgggttatcca
21781 gtggcggtt ttgaagatc tgaaactgtt gtcgttcc ttgggttataa tggcgatc

//

Save the above report in format

Other Formats:

Links:

[LOCUS](#)

[ECORA](#)

NCBI Entrez **Nucleotide QUERY** **BLAST** **Entrez** **?**
Other Formats: **FASTA** **Graphic**Links: **MEDLINE** **Protein** **Related Sequences**

LOCUS ECORAF 5284 bp DNA , BCT 15-JUN-1990
DEFINITION *E.coli rafA, rafB, and rafD genes encoding alpha-D-galactosidase, raf-permease, and raf-invertase, complete cds.*
ACCESSION M27273
NID g147504
KEYWORDS alpha-D-galactosidase; raf-invertase; raf-permease; rafA gene; rafB gene; rafD gene.
SOURCE *E.coli* (strain K-12) DNA.
ORGANISM *Escherichia coli*
Eubacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae; Escherichia.
REFERENCE 1 (bases 1 to 5284)
AUTHORS Aslanidis,C., Schmid,K. and Schmitt,R.
TITLE Nucleotide sequences and operon structure of plasmid-borne genes mediating uptake and utilization of raffinose in *Escherichia coli*
JOURNAL *J. Bacteriol.* 171, 6753-6763 (1989)
MEDLINE 90078124
COMMENT Draft entry and computer readable copy of sequence [1] kindly provided by R.Schmitt, 01-SEP-1989.
FEATURES Location/Qualifiers
source 1..5284
 /organism="Escherichia coli"
 /db_xref="taxon:562"
-35_signal 12..17
-10_signal 36..41
RBS 58..63
 /note="S.D. sequence (ribosome binding site)"
CDS 70..2196
 /note="alpha-D-galactosidase"
 /codon_start=1
 /transl_table=11
 /db_xref="PID:g147505"
 /translation="MISKYCRLLSSPRSDLIKTHPHAEIIWWGSALKHFSPDDCASLERPVANGRLIDITPLTLIAENALGLFSSPGLEGHRNGLDASPVFTVDVEHTENTLRLTSEDSVAGLRLVSELVMTPSGILKVRHALTNLREGDWQINRAITLPVAERAEEVMAFHGRWTREFQPHVRVLTHDAFVLENRRGRTSHEHFPAIIVGTPGFSEQQGEVWAVHLGWSGNHRMRCEAKTDGRRYVQAEALWMPGEKALRKNETLYTPWLYACHSADGLNGMSQQYHRFLRDEIIRFPEQKLRPVLNLTWEGIYFNHNPDYIMQMAERAALGVERFIIDDGWFKGRNDDRAALGDWYTDEQKYPNGLMPVINVHKVSLGMFEGIWEPEMINPDSDLFRLHPDWILSMPGSQPTGRYQYVLNLNIPEAFDYIYKRLWLLGEHPVDYVKWDMNRELVQAGHEGRAAADAQTRQFYRLLDLLRERFPHVEFESCAASGGGRIDFEVLKRTHRFWASNNDAALERCTIQRGMSYFFPPEVMGAHIGHRRCHATFRQHSIAFRGLTALFGHMGLELPVAADAKESDGYRRYALLYKEWRQLIHTGVLWRVDMPDSSIQVQGVVSPDQSQALFMISQLAMPDYTLPGILRFPGLAAEVRYRLRVIDHPEIQLVGEGGHTMRRLPAWMNQPLEASGEWLAKGGIQLPVLDPEASILIALERAV"
RBS 2244..2249
 /note="S.D. sequence (ribosome binding site)"
CDS 2259..3536
 /note="raf-permease"
 /codon_start=1
 /transl_table=11
 /db_xref="PID:g147506"
 /translation="MNSASTHKNTDFWIFGLFFFLYFFIMATCFPFLPVWLSDVVGLSKTDTGIVFSCLSLFAISFQPLLGVISDRLGLKKNLWISISSLVFFAPFFFLYVFAPLLHLNIWAGALTGGVFIGFVFSAGAGAIEAYIERVSRSSGFYEGKARMFGCLGWALCATMAGILFNVDPSSLVFWMGSGGGALLLLLLLARPSTSQTAMVMNALGANSSLISTRMVFLSLFRMRQMWMFVLYTIGVACVYDVFDQQFAIFFRSFFDTPOAGIKAFGFATTAGEICNAIIMFCTPWIINRIGAKNTLLVAGGIMTIRITGSAFATTMTEVVLKMLHALEVPFLLV"

GAFKYITGVFDTRLSATVYLIGFQFSKQLAAILLSTFAGHLYDRMGFQNTYFVLGMIV
 LTVTVISAFTLSSSPGIVHPSVEKPAHSEIN"
 RBS 3523..3528
 /note="S.D. sequence (ribosome binding site)"
 CDS 3536..4966
 /note="raf-invertase"
 /codon_start=1
 /transl_table=11
 /db_xref="PID:g147507"
 /translation="MKQRLSLAQSALEKLSARRGNTWYPIFHLAPPAGWMNDPNGLIY
 FNGRYHAFFQHHPASAYQGPMHWGHATSTDMHWQHELVAPGDKYDRDGCFSGSAV
 DDDGVLISLIYTGHICLEDRGNDSIIREVQCLATSHDGIRFEKQGCVLTPPEGIMHFRD
 PKVWHEDGSWMVIGARDASDNGQVLLYRGTSLRDWHLHVAHSAAGESYMWECPDF
 FRCGNFHWMFSPQGMNPSPYRFRNLQSGVLAGNWKPGSVFALKGVFEELDYGHDFY
 APQSMILAEDGRRIIMAWNMNWMDSPVPTRSEAWAGCLTLPREVDGRLCQRPVREVE
 SLRRKCQPLSPVRLHGVQLTENVQAAELLVTWHTVDHAEHYGIRLGEGLRFYVDNQ
 AGRLILWRYYPEEGLDGYRSVELPDTEYLTLRIFLDRSSVEVFVNDGEATLSSRIYPQ
 ADSRQLSLYAAHGDAILTDGTLWMLT"

BASE COUNT 1199 a 1166 c 1406 g 1513 t

ORIGIN

1 aaacgttttg gttgatgttc gaaacgtttc ggatcaacag taagacatac ctgaaagcgg
 61 agatgtctta tgatttcaaa gtactgcaga ctgagcagtc ctcgctctga ttttaattatt
 121 aaaacccatc cgcgcgcaga aattatctgg tggggctctg cactgaaaca tttctcaccg
 181 gatgactgtg ccagcctgga aagaccagg gcgaatggc gtctggatat tgatacgcca
 241 ctgacactga tcgctgaaaa tgcgccttgcgc cttttagtt ctccggact ggaaggacac
 301 agaaatggac tggatgcattc tcctgtttt tatacagttt acgtggaaaca taccgaaaac
 361 accctgagac ttaccagtga agattcggtt gccggcctgc gtctggtcg cgagctgg
 421 atgacgcccattt cggggattctt gaaagttcgat catgcactga ccaacctcag agagggagac
 481 tggcagataat atcgtttcgc aatcacttta cctgttagctt aacgtgcggg agaagtcatg
 541 gccttcacgc gacgcgtggac tcgtgaattt cagccgcaca ggttacgtt tactcatgt
 601 gcttttgcgc tggaaaatcg cagaggggcg acatctcatg agcattttcc ggcgctgatt
 661 gtcggcacac caggcctctc gaaacaacag ggagggatgtt gggctgtgc tctgggtgg
 721 atggaaatcc accgcatttgc atgtgaggca aaaactgttgc gcaaggcgtt cgtacaggct
 781 gaggctctgtt ggttgcgggg tgagaaggctt ctcaggaaga atgaaaccctt gtacaccccg
 841 tggctatattt cctgcacttgc tgcggatggc ctgaatggaa tgagtcaatc ataccatcg
 901 tttttgcgtt atgaaattat cctgttccctt gggcccccgtt acatctcaat
 961 acctggaaatcc ttattttttt caatcacaat cctgttccctt tcatgcagat ggctgagcgt
 1021 gcagcagcac tggcgttgc acgtttcattt attgtatgtt gctggttttaa aggacgttac
 1081 gatgaccgcgc cggcctctggc cgaatggat accgttgcac agaaataccctt gaaacggctg
 1141 atgcccgtttaa ttaatcatgtt gaaatcttc ggtatggatgtt ttgttatatg gtttgaacca
 1201 gaaatgatattt atccggattt tgacctgtttt cgttccatc cggactgtt atttcaatg
 1261 ccaggatattt ctcagccaaatcc cggaaatggat cagtatgtt ttaacctgaa tattccggag
 1321 gccttgcattt atatttataa acgtttcttgc tggtaactt gagaacatcc ggttggatt
 1381 gtggaaatggg acatgaatcg tgagctgttgc caggcaggccat gtaaggccg tgccgcgc
 1441 gatgcacaga cccgttgcgtt ctatcgatgg cttgtatctcc tccgttgcacg tttccacat
 1501 gttgagtttgc agtccgttgc ttccgggtttt gggcgatgtt acgttgcgtt cctgaaacgc
 1561 acacaccggcgt tctggcacttgc tggatgttgc acgttgcac catacaacgt
 1621 ggcatgatgtt actttttccc ttcttgcgttgc atgggggcac atattggccat tcggcgttgc
 1681 catgcaactt tccggcagca cagcatcgat tttcgatggc tgacggcattt gttccgcatt
 1741 atggggctgg agtggatcc ggttgcgc gatgcgttgc aatctgttgc ttatcgccgg
 1801 tatgccttgc tctataaaga atggcgacaa ctgatttgcata caggtgttgc tggcgttgc
 1861 gatgttgcgg attcttcgttgc acgttgcgttgc ggatgttgc gcccgttgc gtcgttgc
 1921 ctttttatgtt tcagccatgtt tgcaatggcc gattacaccc taccaggcat acttcgttt
 1981 cccggactgg cggcagaatgtt ggttgcgc gatgcgttgc ttatcgccgg gaaattcaat
 2041 ctgggttgcgtt aagggttgcgttgc taccatgttgc acgttgcac cctggatgaa ccagccctc
 2101 gaggccatgtt gtggatggctt ggttgcgttgc ggatgttgc gcccgttgc gtcgttgc
 2161 agcgcgttgcgttgc tggatgttgc acgttgcgttgc ggatgttgc gcccgttgc gtcgttgc
 2221 gtcctgttttcaatattt cccgttgcgttgc tggatgttgc gatgcgttgc ttatcgccgg
 2281 aaaatactgttgc tttcttgcgttgc tggatgttgc acgttgcac cctggatgaa ccagccctc
 2341 cctgtttccgttgcgttgc tggatgttgc acgttgcgttgc ggatgttgc gcccgttgc gtcgttgc
 2401 caggatgttgcgttgc tggatgttgc acgttgcgttgc ggatgttgc gcccgttgc gtcgttgc
 2461 tcatatcaga tccgttgcgttgc tggatgttgc acgttgcgttgc ggatgttgc gtcgttgc
 2521 tatttttgcgttgc cccgttgcgttgc tggatgttgc acgttgcac cctggatgaa ccagccctc
 2581 cagggttgcgttgc tggatgttgc acgttgcgttgc ggatgttgc gcccgttgc gtcgttgc
 2641 ttgttgcgttgc tggatgttgc acgttgcgttgc ggatgttgc gcccgttgc gtcgttgc

2701 tgttcgggtg tctgggttgg gcgttatgtg caactatggc cgaaatactt tttaatgtcg
2761 atccctctct ggtttctgg atggggtcag gaggcgcatt attgttgcgtg cttctgttgt
2821 atctggcgcg ccccagtacc agccagacgg caatggttat gaatgcactg ggtgcatt
2881 ctccctgat ttctaccaga atggtctca gcctgttgc tatgcgtcag atgtggatgt
2941 ttgttctata tacgataagg tggcctgtg tctatgtatgt atttgatcag cagttgcca
3001 tatttttcg ttcattttt gacactccgc aggcaagaa aaaggcattc ggatttgc
3061 ccaactgcggg ggagattgt aatgccatta tcatgttctg tacaccatgg ataattaatc
3121 gcattgggtgc caaaaataacc ctgcttgcg cgggggaat tatgactatc cgcattaccg
3181 gttctgctt tgccaccacc atgacagaag tggtattct gaaaatgtt cacgcttgc
3241 aagttccatt tttgctgggtt gggcattca aatatattac gggcgtgtt gacacccgac
3301 tgcagcgtac ttttattta attgggtttc agtttccaa acaactgtt gcaataactc
3361 tctctaccc tggccggcac ctgtatgatc gtatggatt ccagaatacg tattttgtgc
3421 tcgggatgat ttttctgact gttaccgtt ttttgcgtt cttttttttt aactgtgaa
3481 ggatcgtaaa cccttctgtt gaaaagccc ctgtacaca ttcggagatt aactgtgaa
3541 acagcgtctt ttttggcac agtctgcctt tgaaaaactt agcgcacgtc gcgtaataac
3601 ctggtacccg attttcattc tggctccacc tgccgggtgg atgaatgatc caaatggcct
3661 gatttatttc aatggacgtt accatgcgtt cttccagcat catccagca ggcacatata
3721 ggggccaatg cactggggc atgcccaccat taccgacatg ttgcactggc aacacgagct
3781 tgcgcgtctg gcaccggag ataaatatga tgcgtatggc tttttttttt gggatggcgt
3841 ggatgtatgat ggcgtctat cacttattta taccggccat attttgtctg aagatcggtt
3901 taatgacagg attttccgtt aagtacatgt tctgttacc tttttttttt atgcattttc
3961 tgagaaggcgg ggcgtgttgc tgacacccccc cgaaggattt atgcattttc
4021 agtctggcac gaagacggct cttttttttt ggttattttt gcccgggacg
4081 tggcaagtt ctgttgcattt gcccggacatc tttttttttt tggcattttt
4141 tgctcattcc gcagccggag aagttatattt gttttttttt
4201 taattttcac tggctgtatgt tttttttttt
4261 taacctttt cagagcggtt tttttttttt
4321 gaaaggggta tttttttttt
4381 ggctgaggac ggcaggcgta tttttttttt
4441 taccctgtatgt gttttttttt
4501 tggccggctg tgccagcgac tttttttttt
4561 attatccccct gtaagggttac tttttttttt
4621 attactgggtt acgtggcata tttttttttt
4681 agaagggtctg cgggttctatg tttttttttt
4741 ccctgaggaa gggctggatg tttttttttt
4801 tcttcgtatt tttttttttt
4861 cttatcaagc cgtatttttcc tttttttttt
4921 tggcgatgcg atattaactt tttttttttt
4981 agtattttttt cctgacatta tttttttttt
5041 tgaaaataat ggagaatgaa tttttttttt
5101 tgattttttt ccggagtcat tttttttttt
5161 gcactggat tttttttttt
5221 gttcaatatt tttttttttt
5281 cgac

//

Save the above report in format
